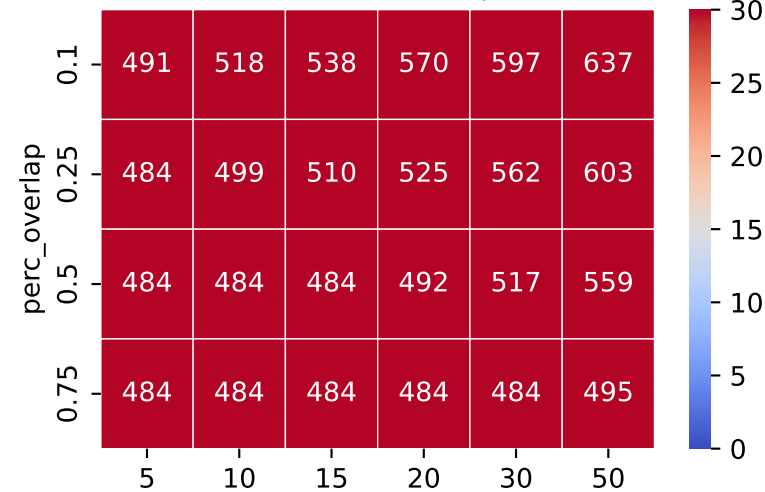


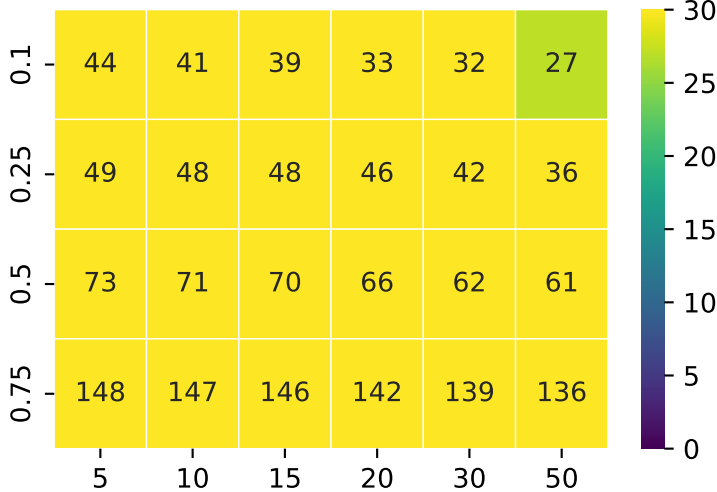
Projection: PCA (n_components=1)

Clustering: DBSCAN (eps=1.0)

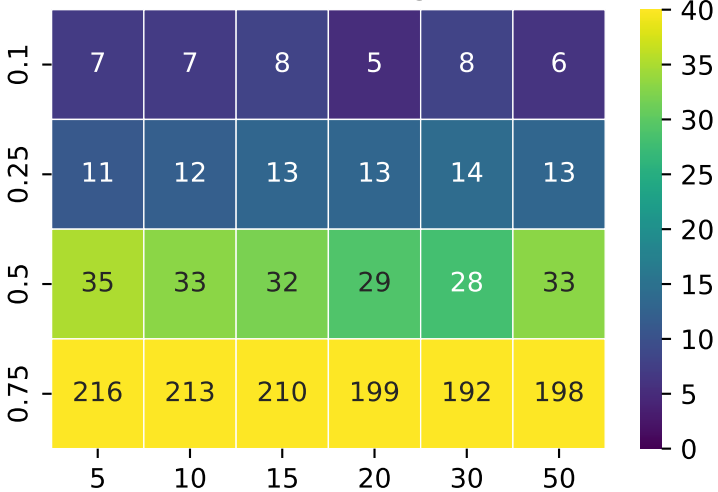
Unclustered samples



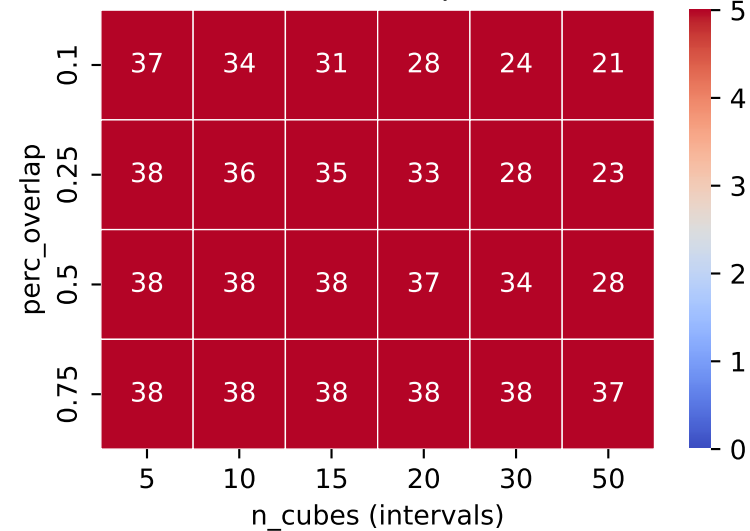
Number of Clusters



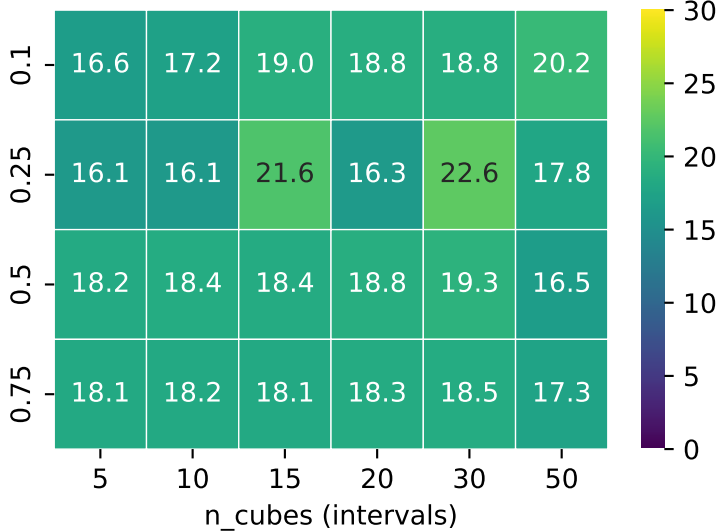
Number of Edges



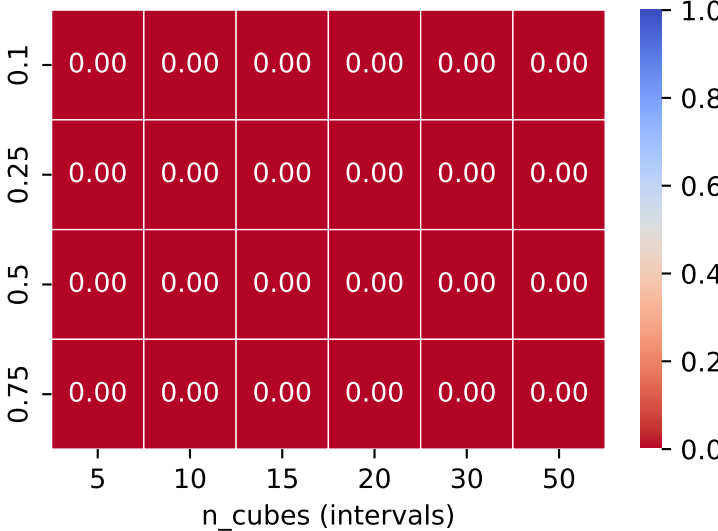
Connected Components



Cluster Size Variation

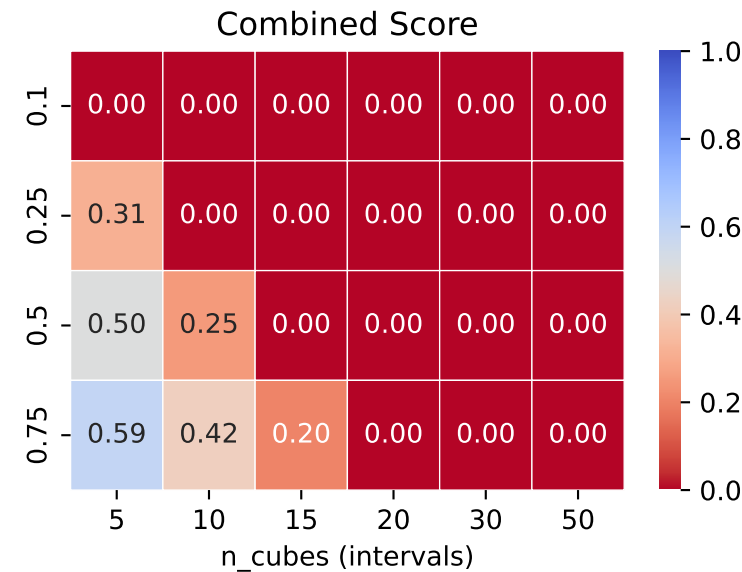
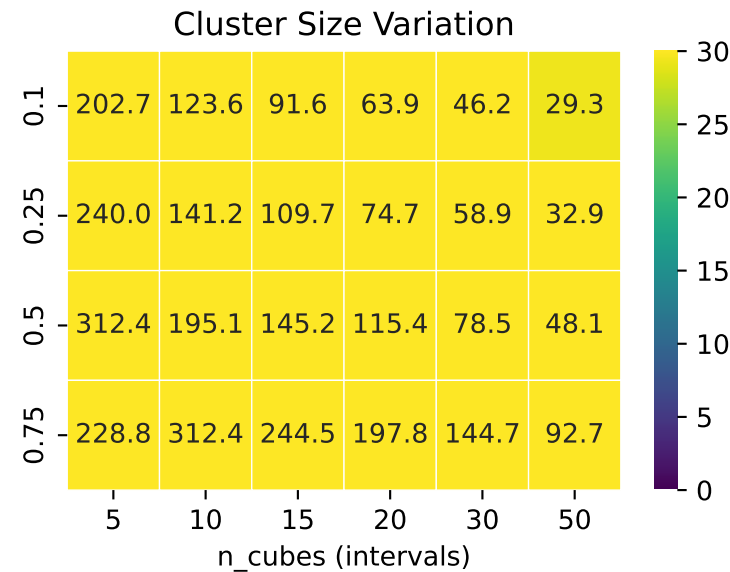
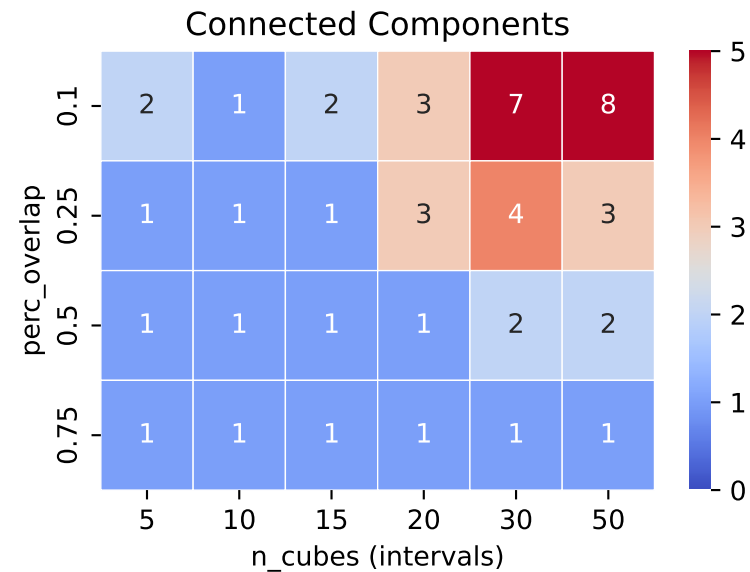
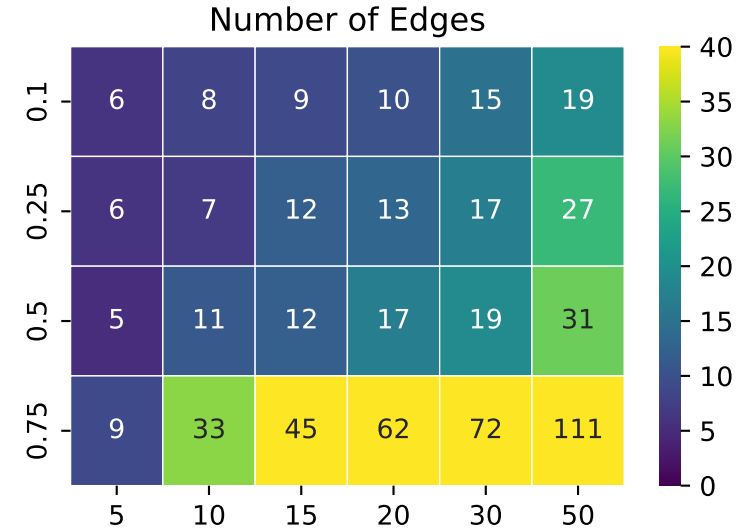
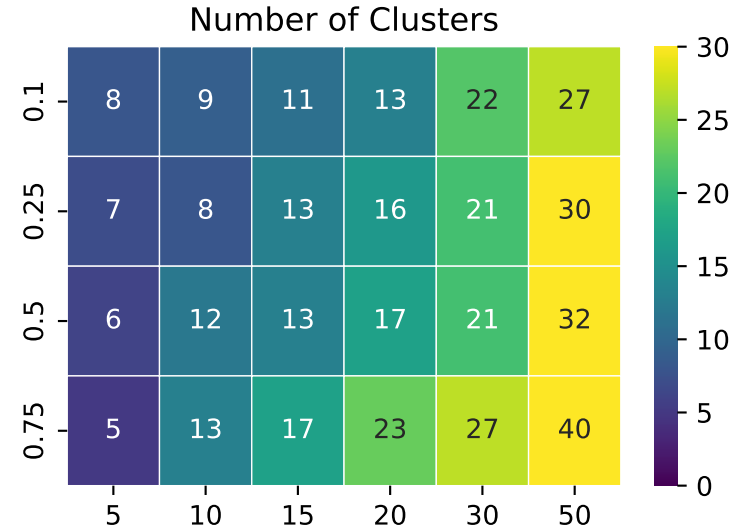
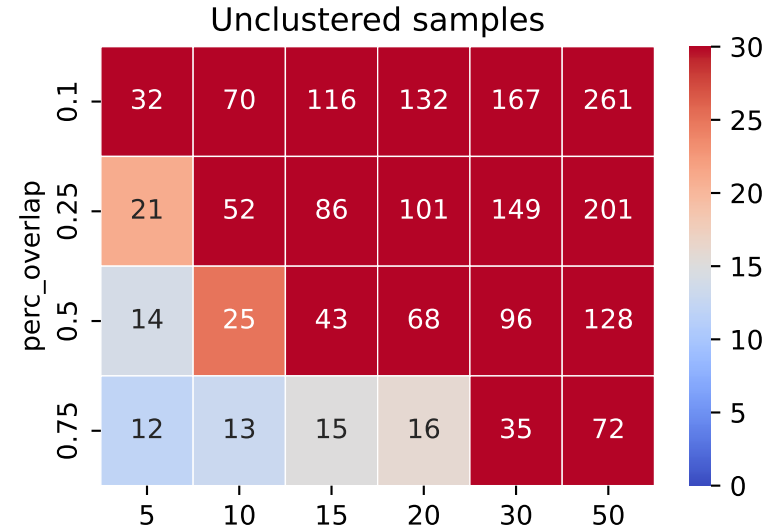


Combined Score



Projection: PCA (n_components=1)

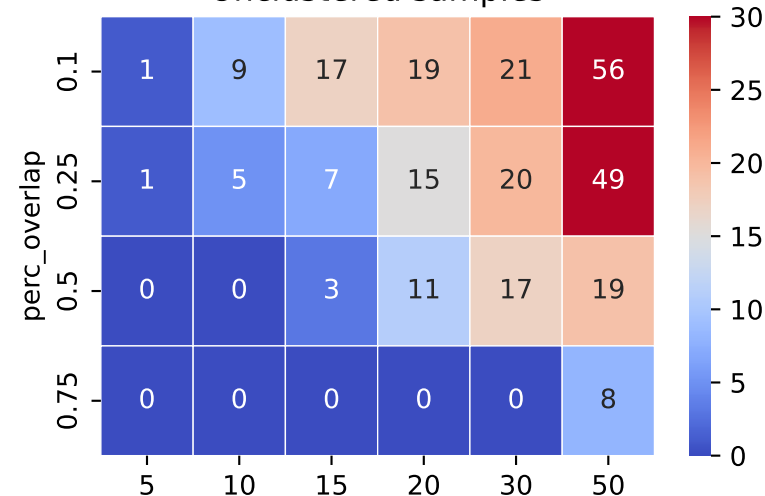
Clustering: DBSCAN (eps=3.0)



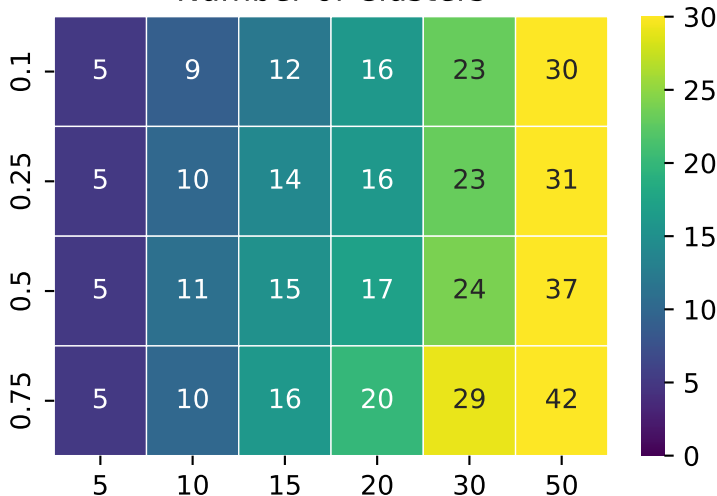
Projection: PCA (n_components=1)

Clustering: DBSCAN (eps=5.0)

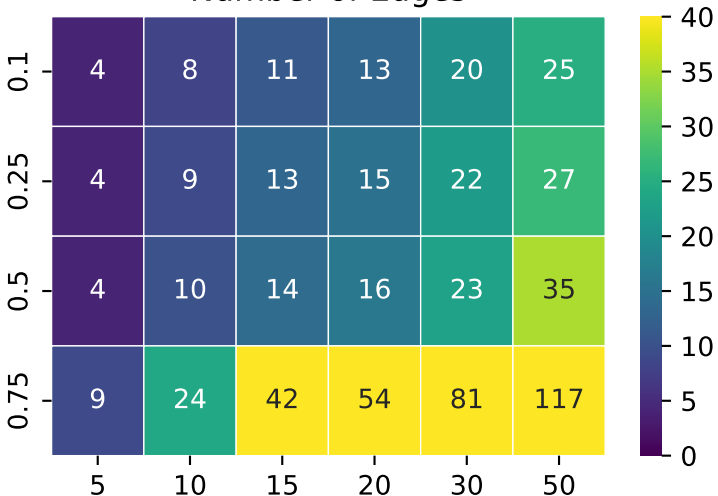
Unclustered samples



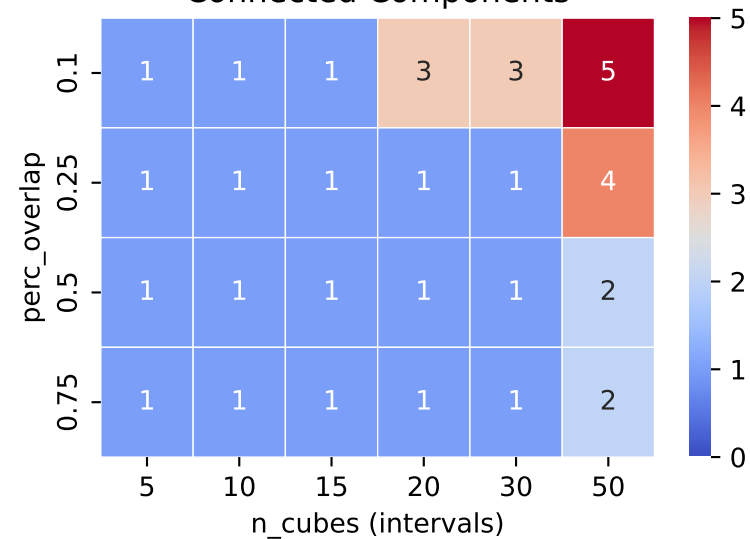
Number of Clusters



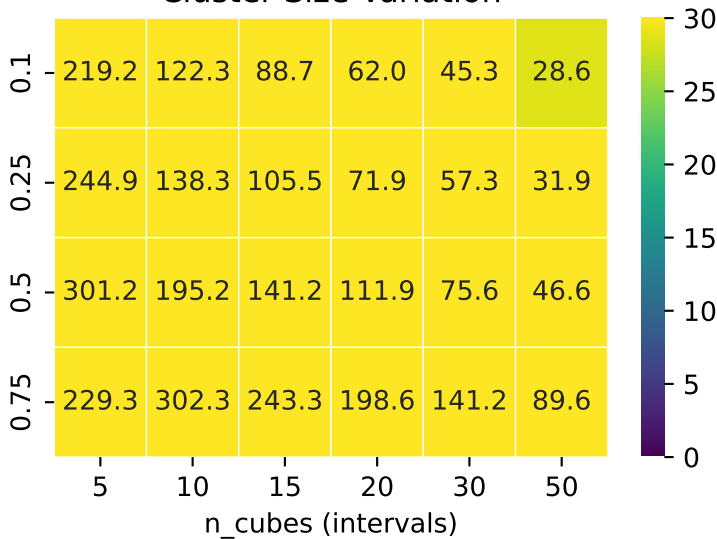
Number of Edges



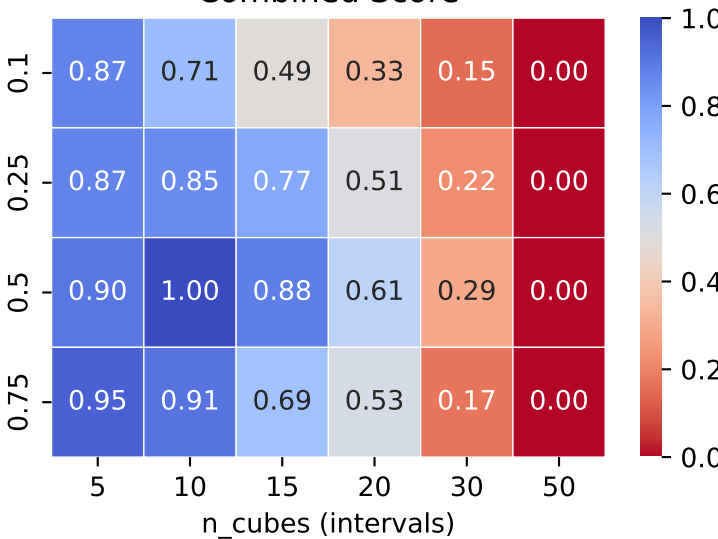
Connected Components



Cluster Size Variation



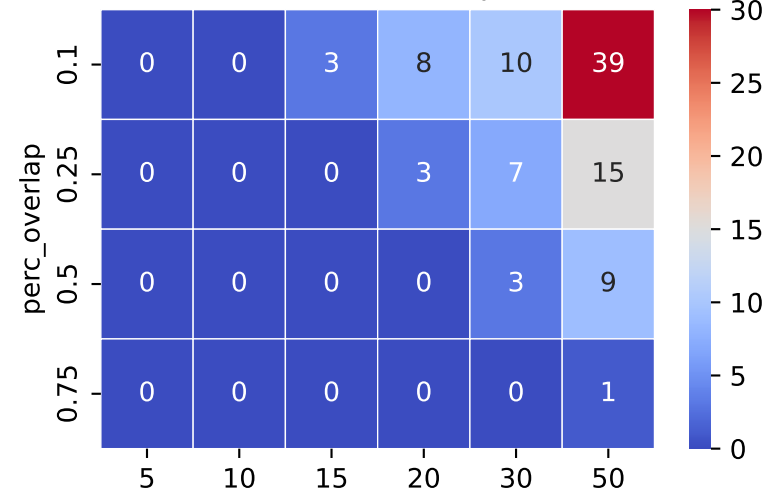
Combined Score



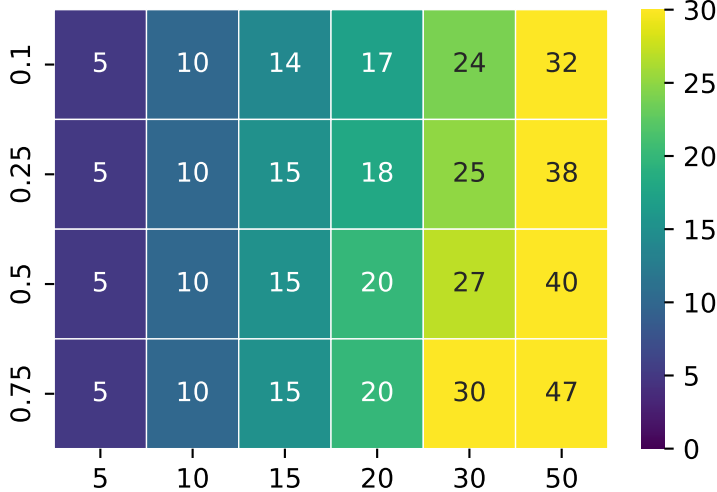
Projection: PCA (n_components=1)

Clustering: DBSCAN (eps=10.0)

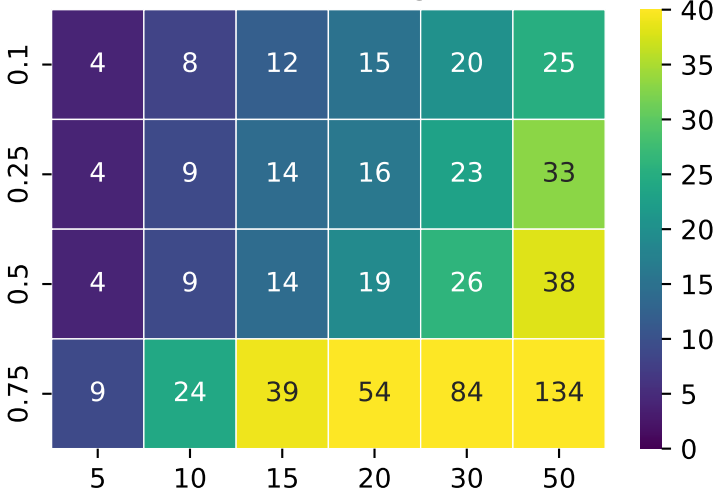
Unclustered samples



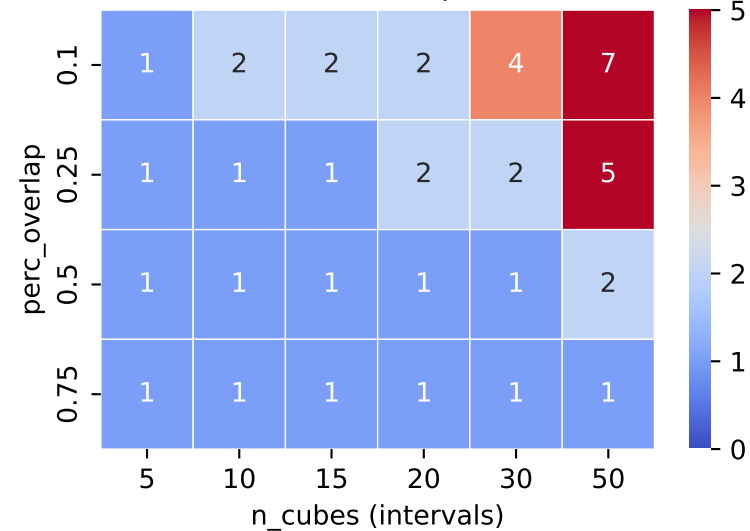
Number of Clusters



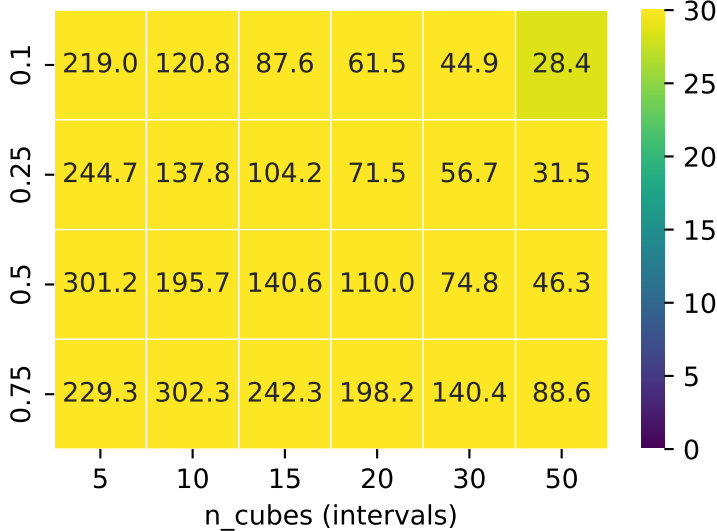
Number of Edges



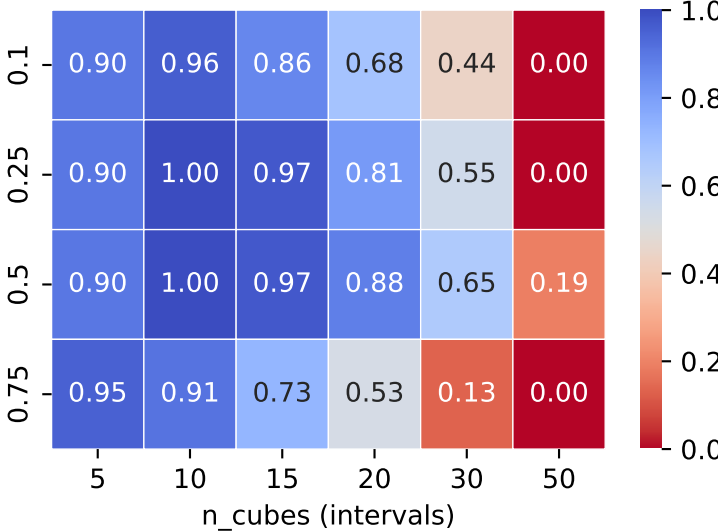
Connected Components



Cluster Size Variation

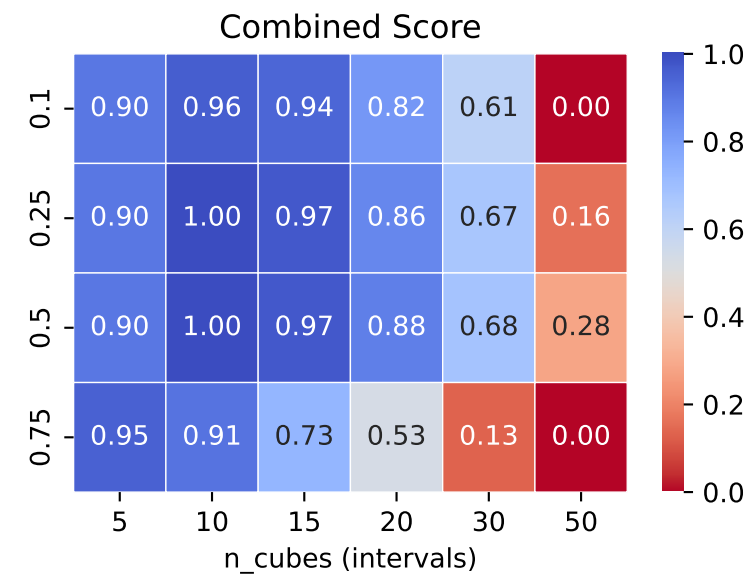
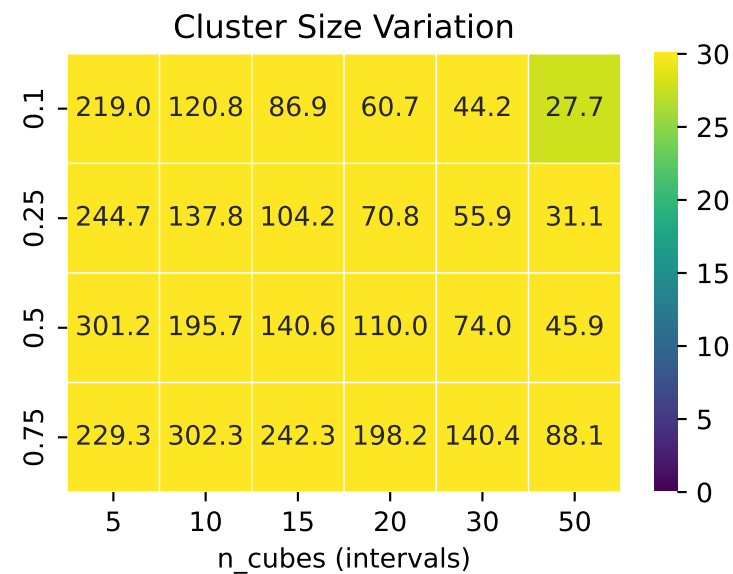
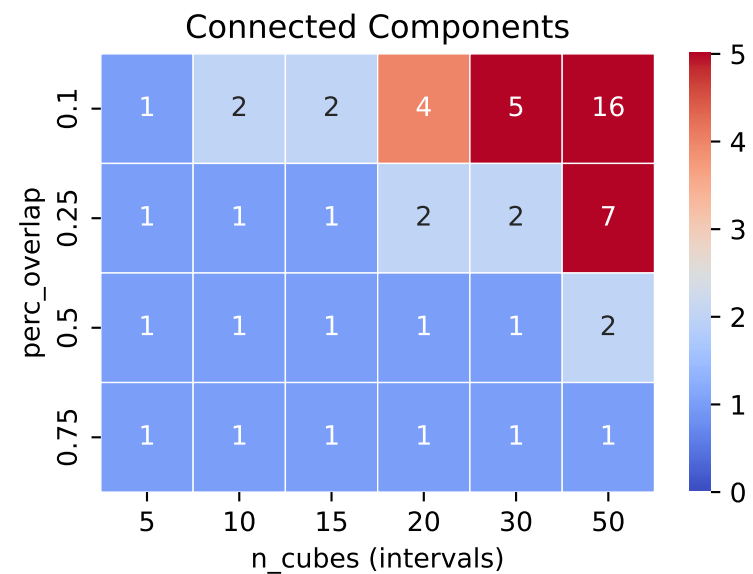
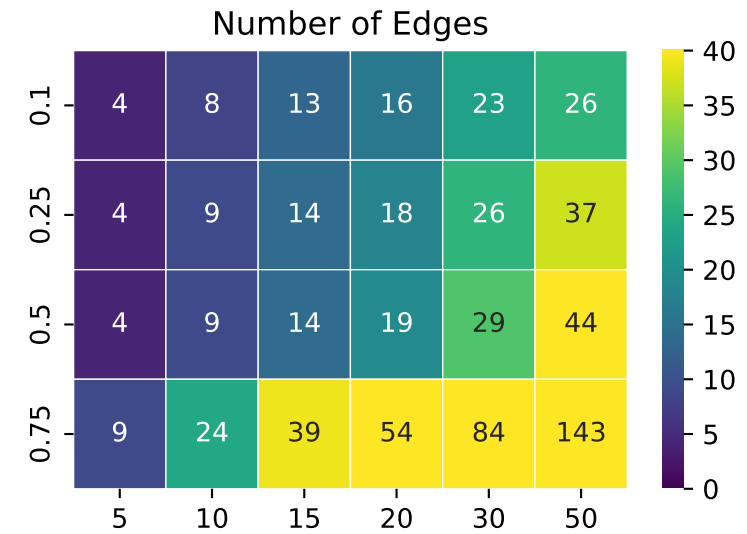
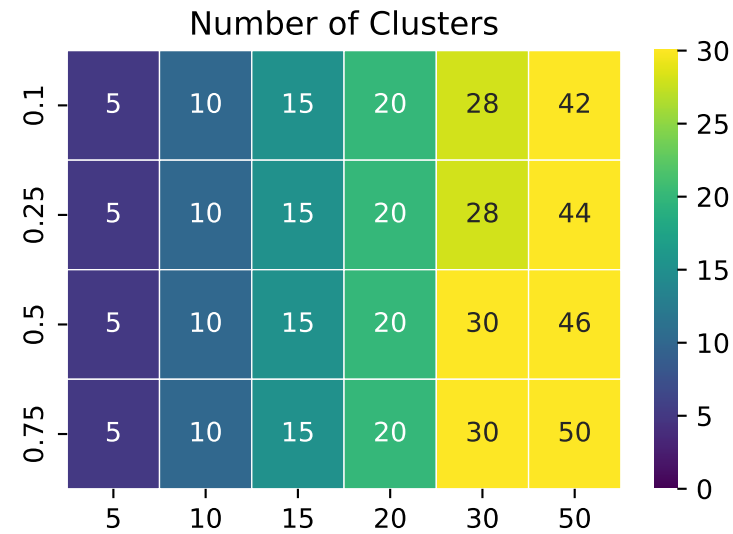
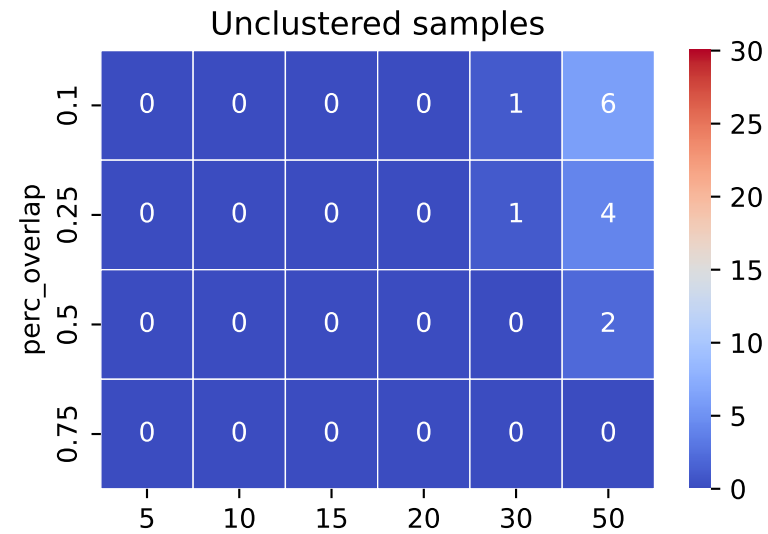


Combined Score



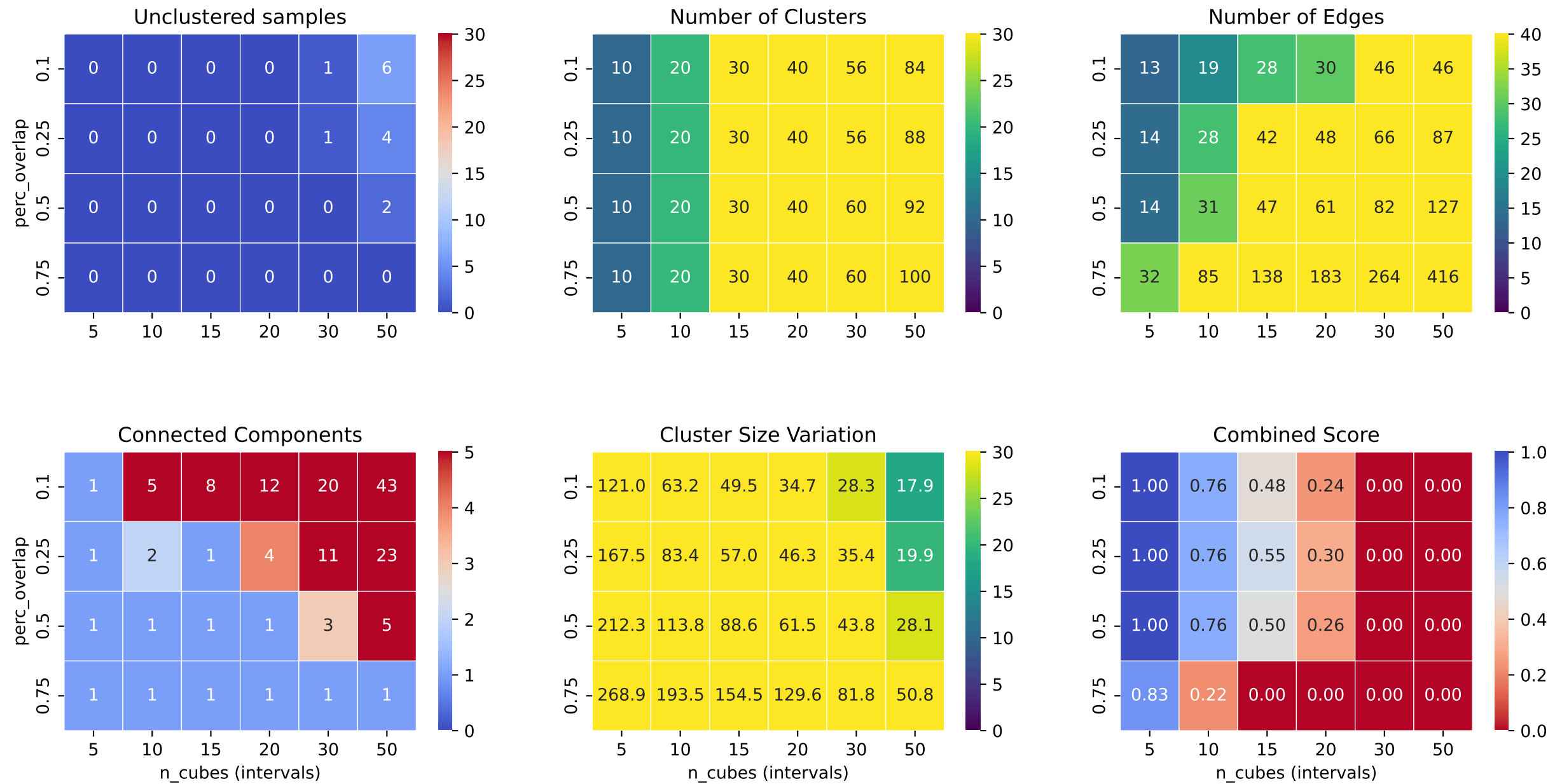
Projection: PCA (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=1)



Projection: PCA (n_components=1)

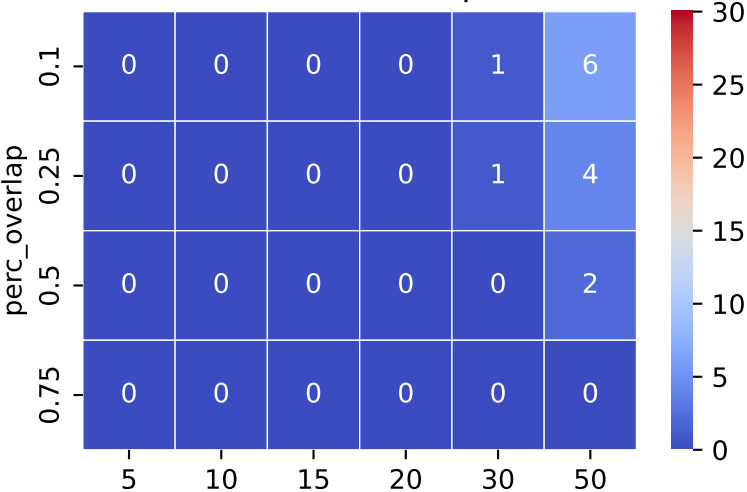
Clustering: Bayesian Gaussian Mixture (n_components=2)



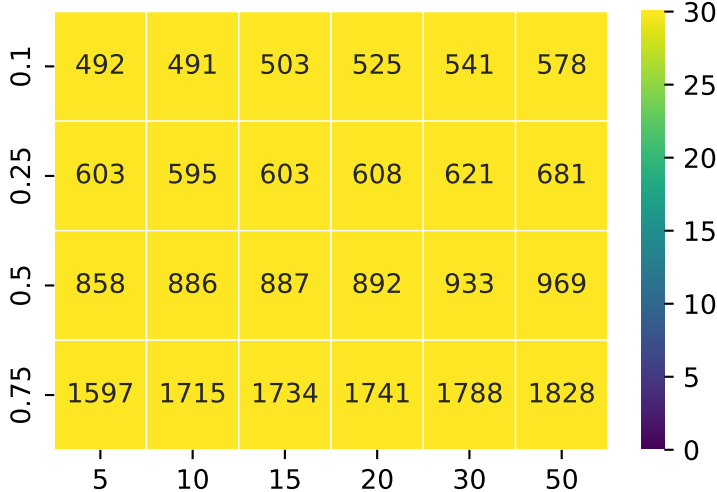
Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=1.0)

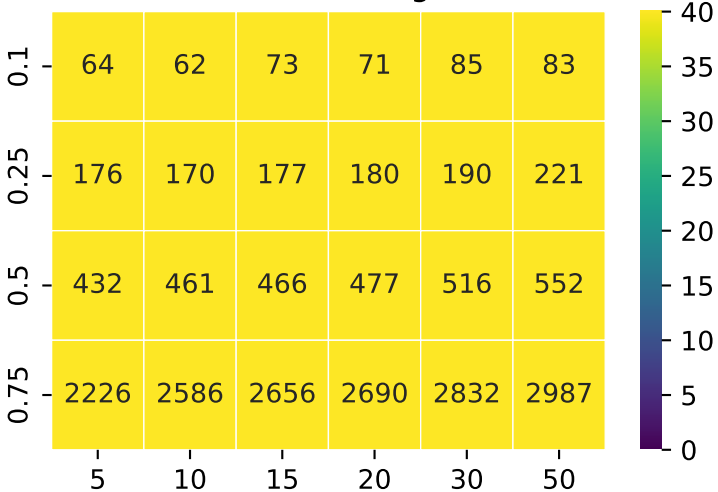
Unclustered samples



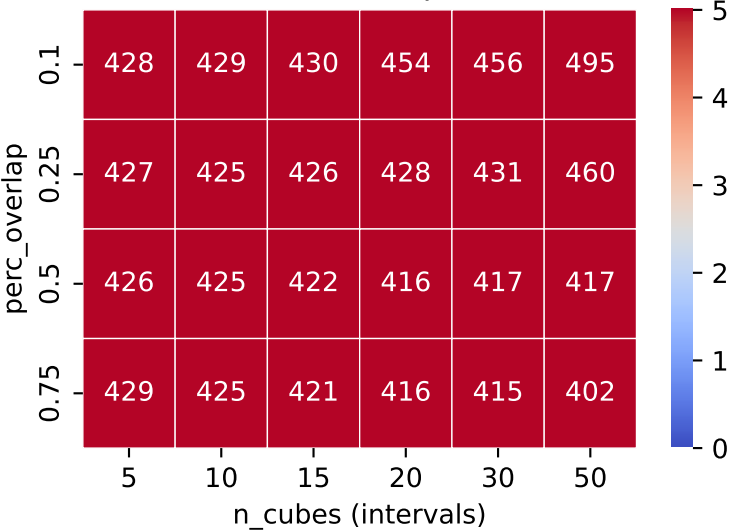
Number of Clusters



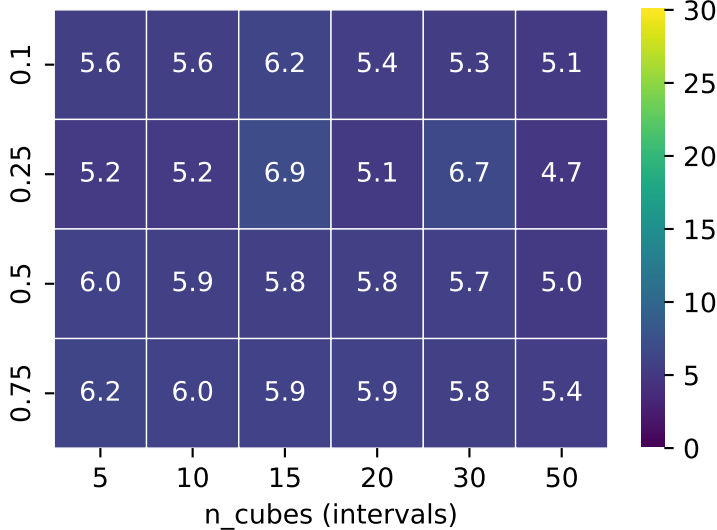
Number of Edges



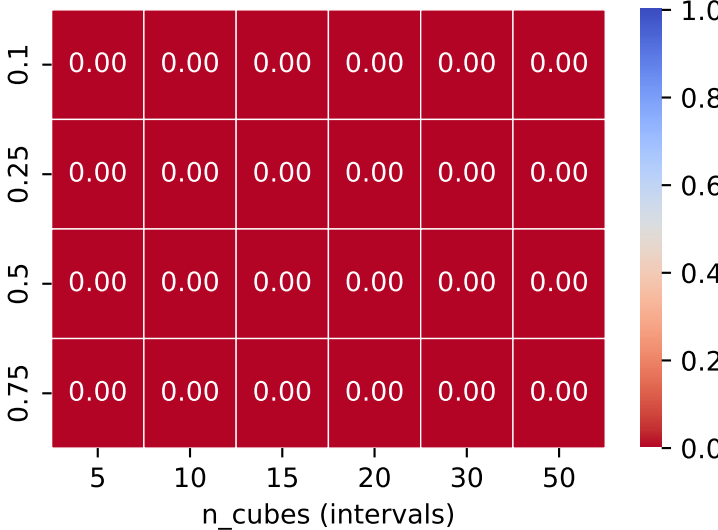
Connected Components



Cluster Size Variation

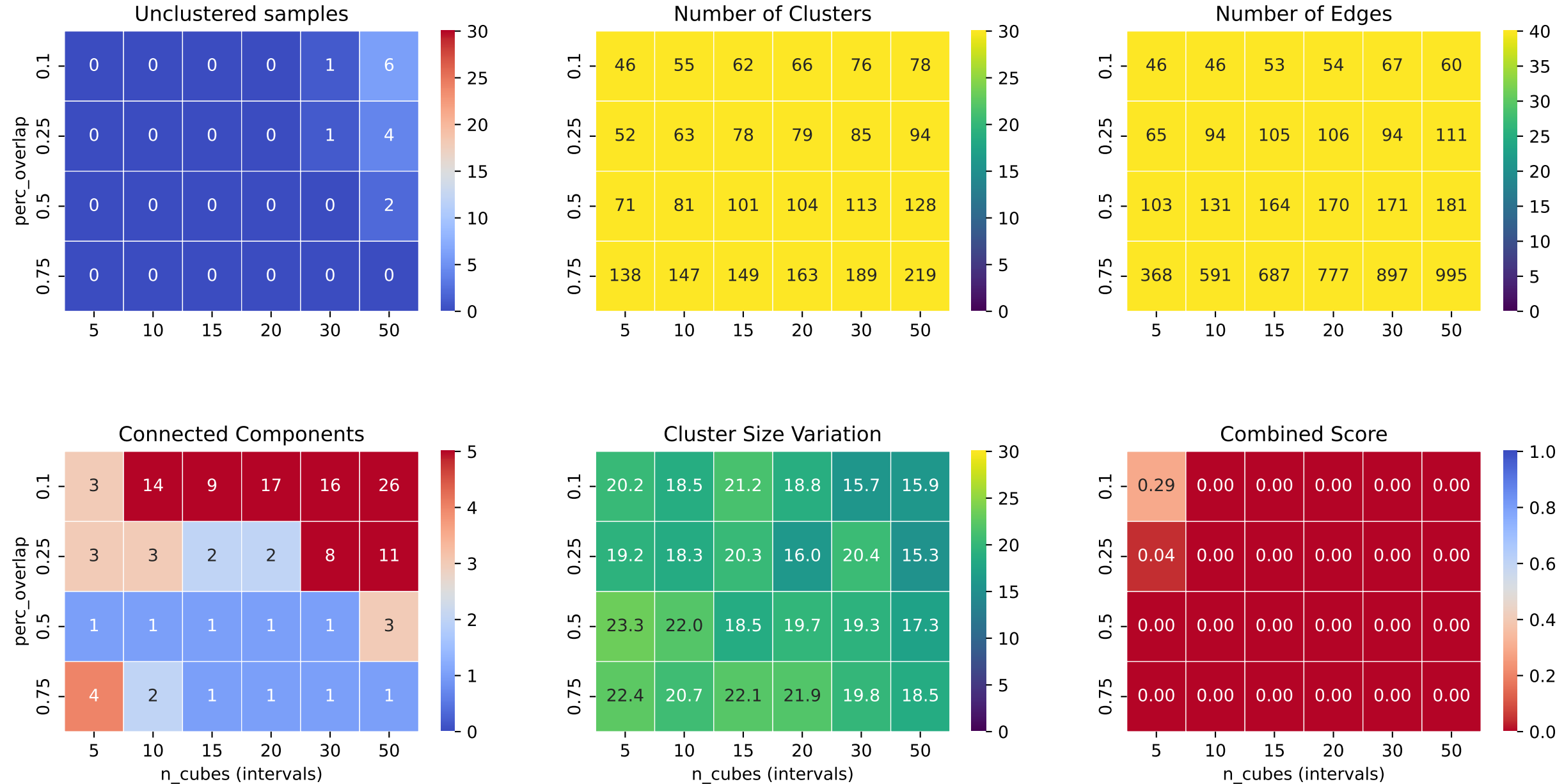


Combined Score



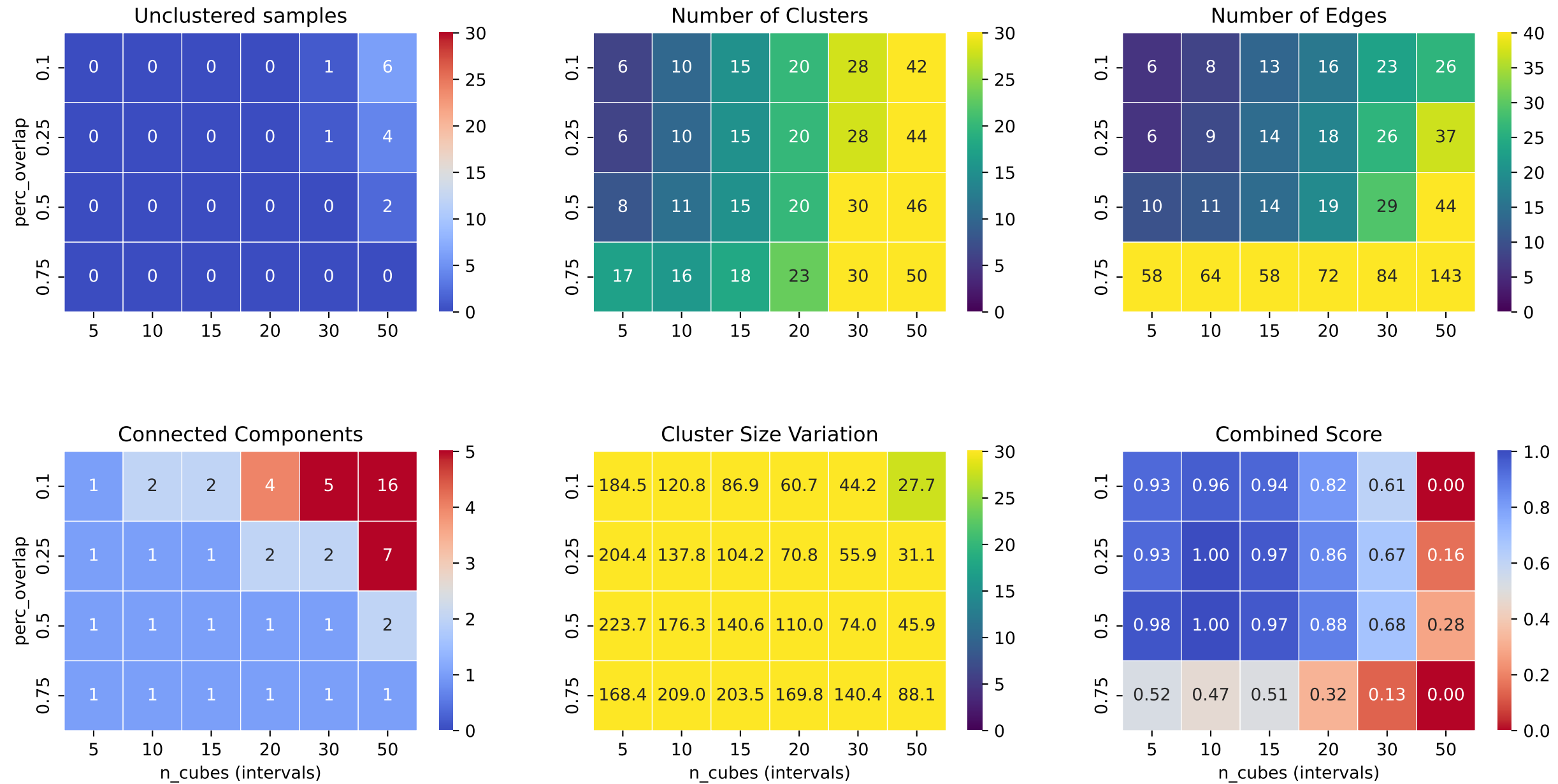
Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=10.0)



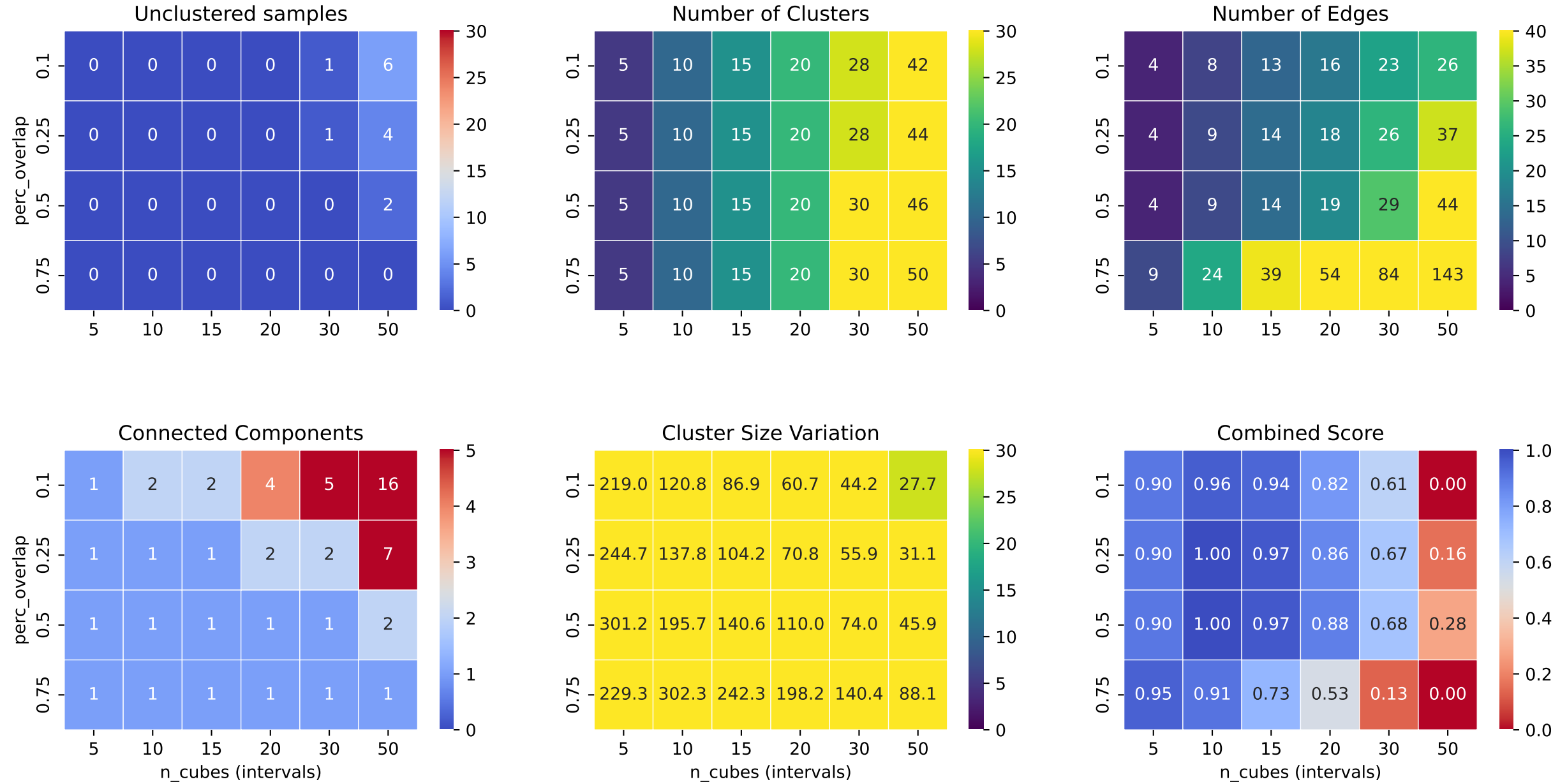
Projection: PCA (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=30.0)



Projection: PCA (n_components=1)

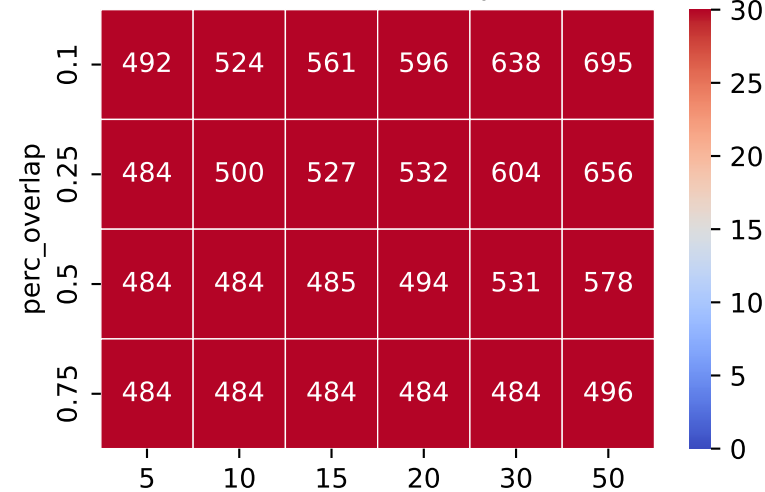
Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)



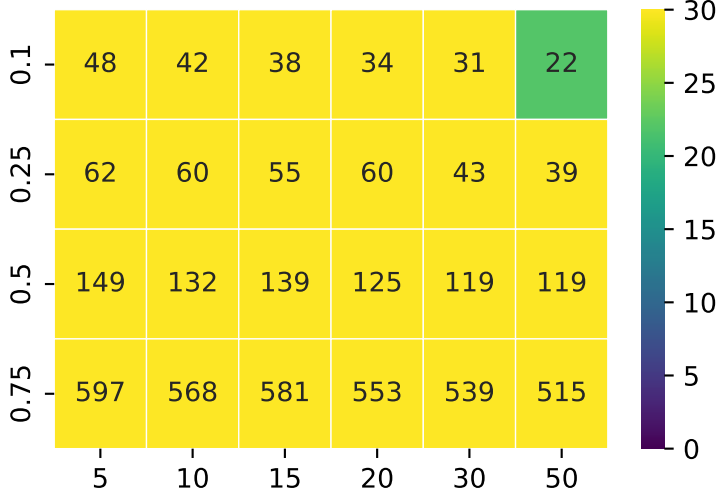
Projection: PCA (n_components=2)

Clustering: DBSCAN (eps=1.0)

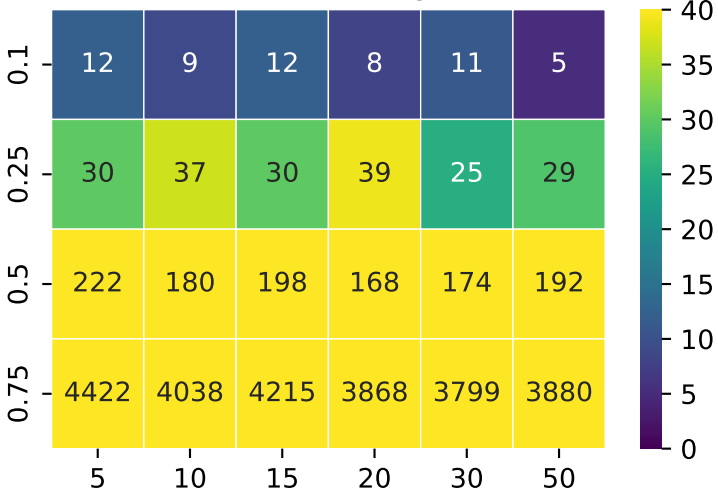
Unclustered samples



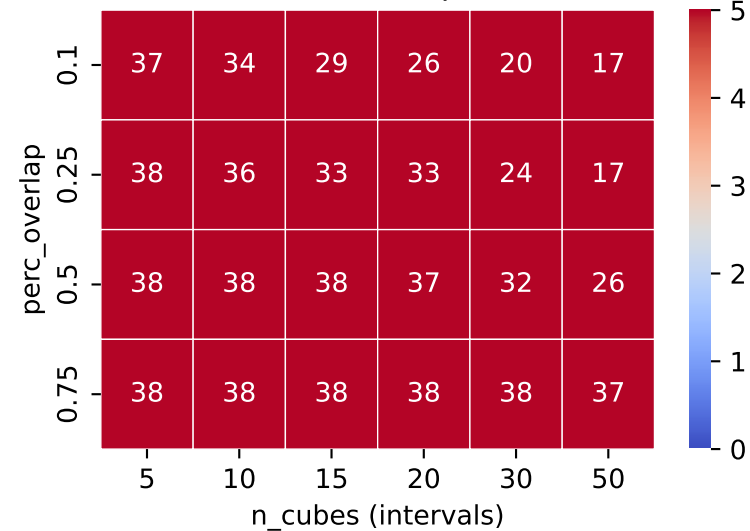
Number of Clusters



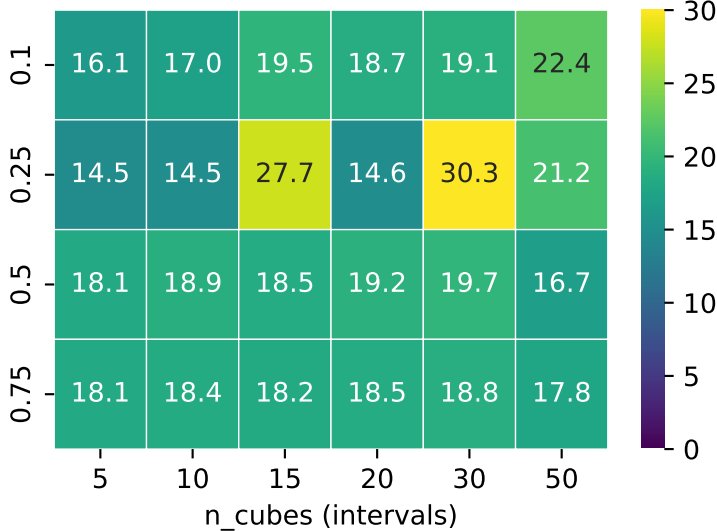
Number of Edges



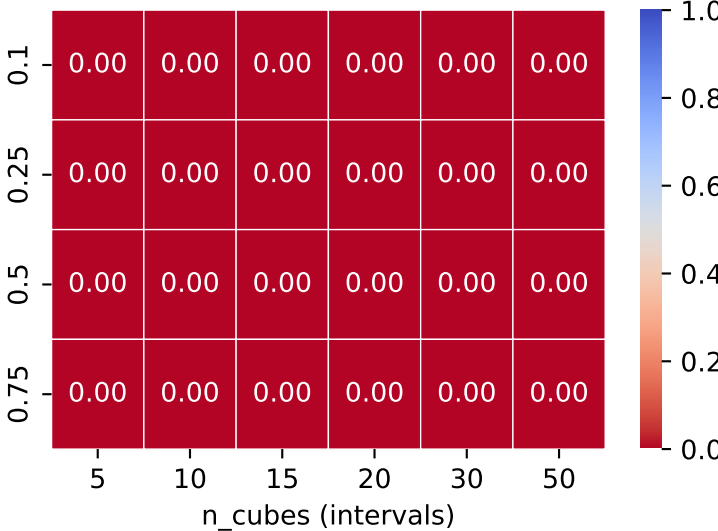
Connected Components



Cluster Size Variation



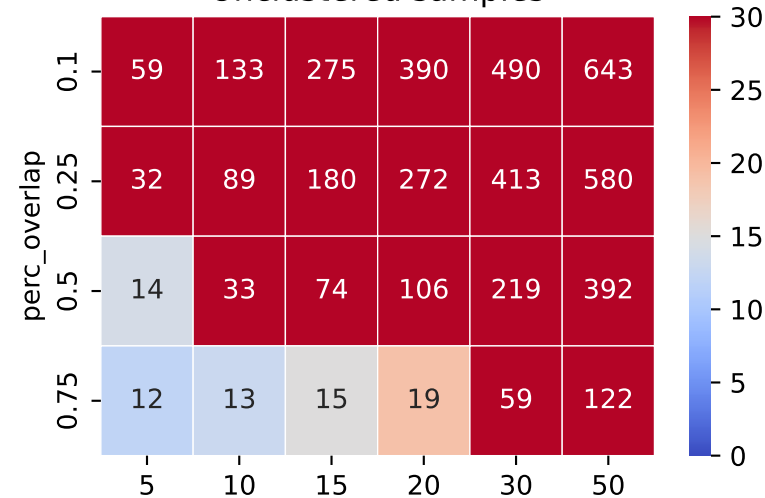
Combined Score



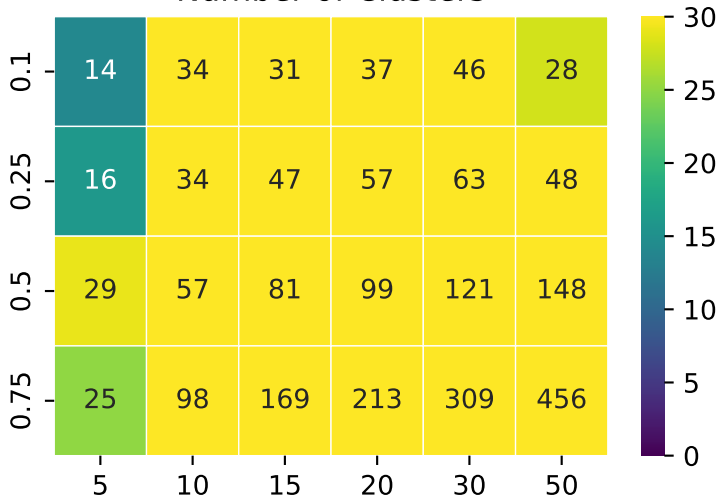
Projection: PCA (n_components=2)

Clustering: DBSCAN (eps=3.0)

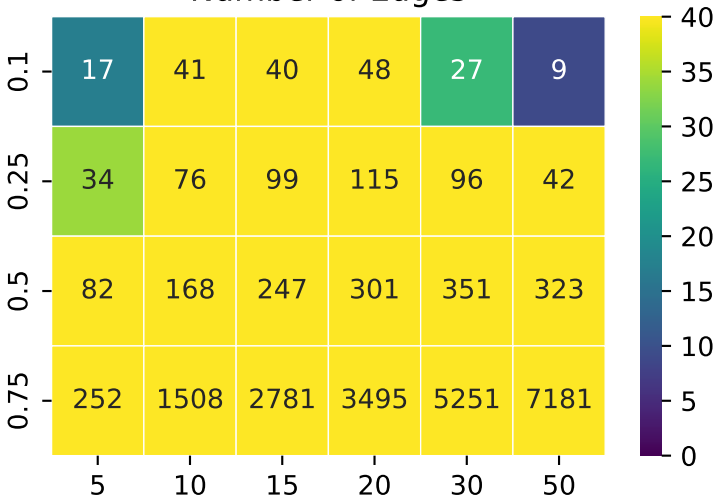
Unclustered samples



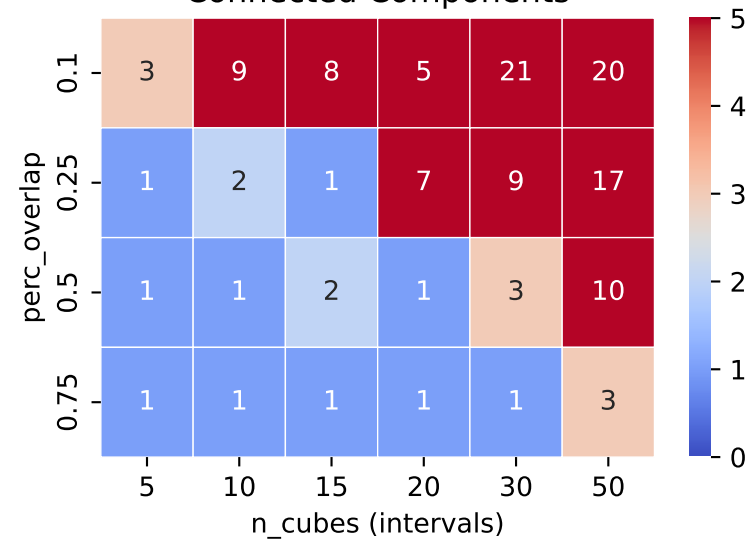
Number of Clusters



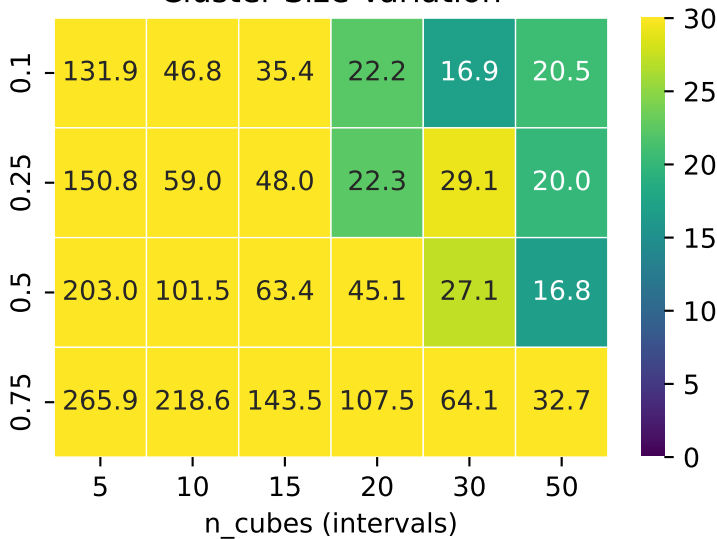
Number of Edges



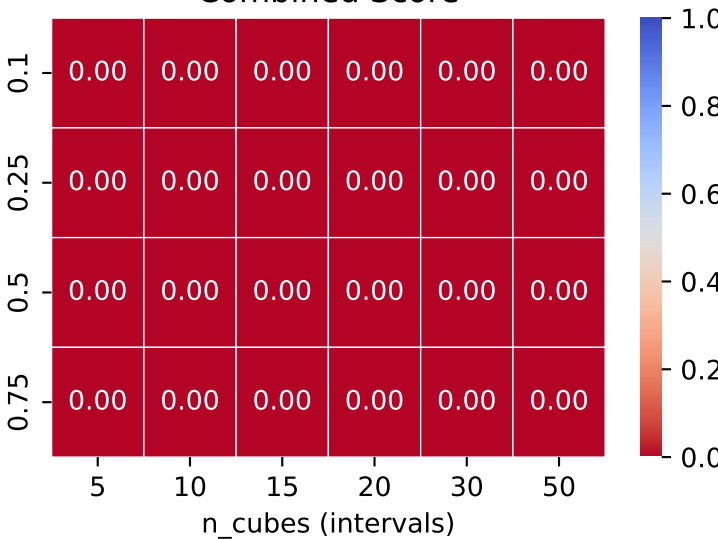
Connected Components



Cluster Size Variation



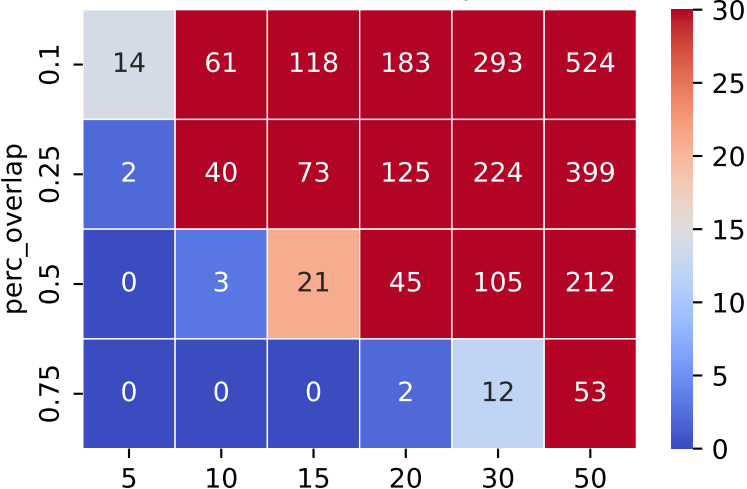
Combined Score



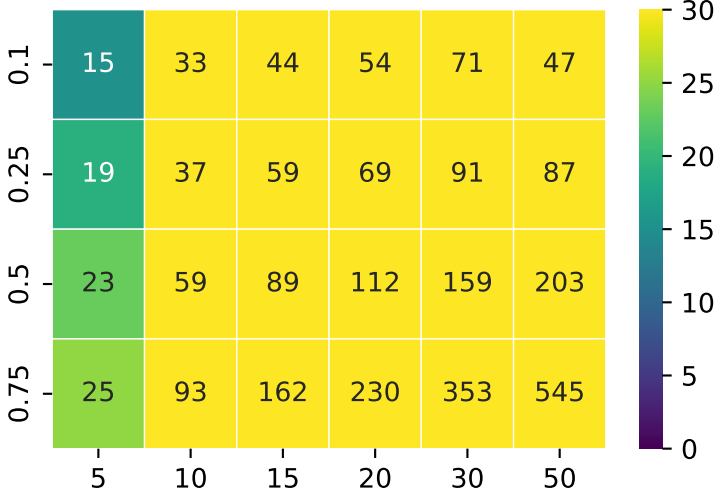
Projection: PCA (n_components=2)

Clustering: DBSCAN (eps=5.0)

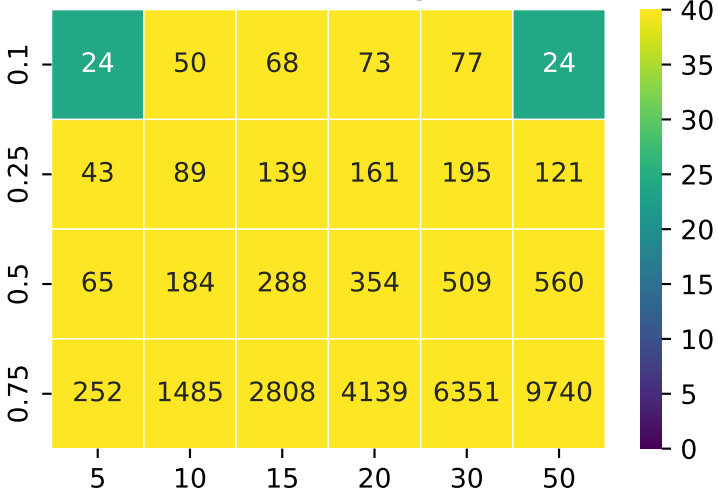
Unclustered samples



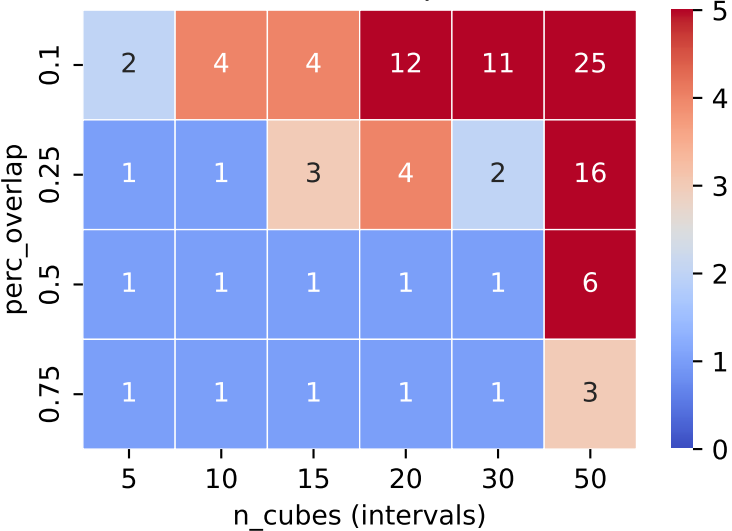
Number of Clusters



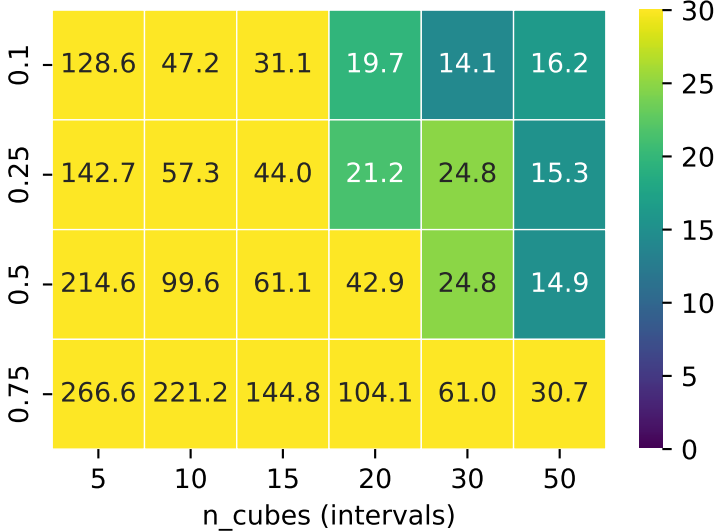
Number of Edges



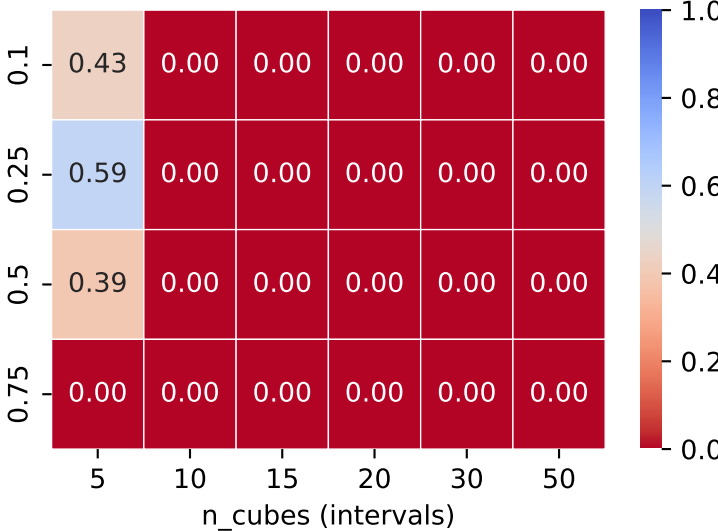
Connected Components



Cluster Size Variation



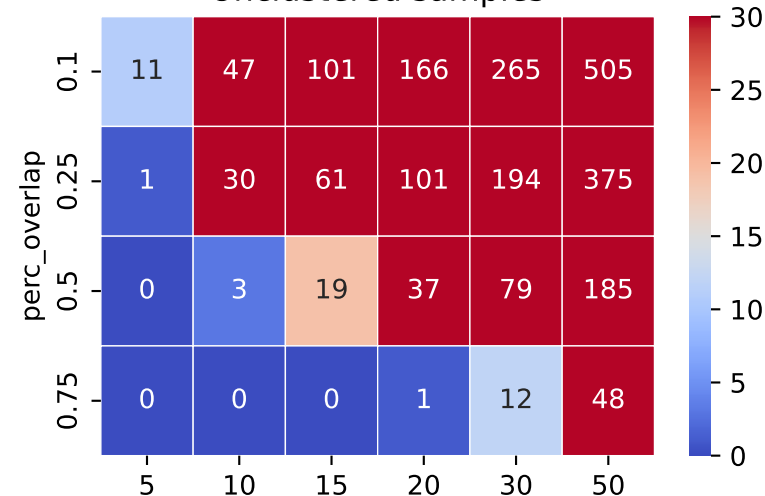
Combined Score



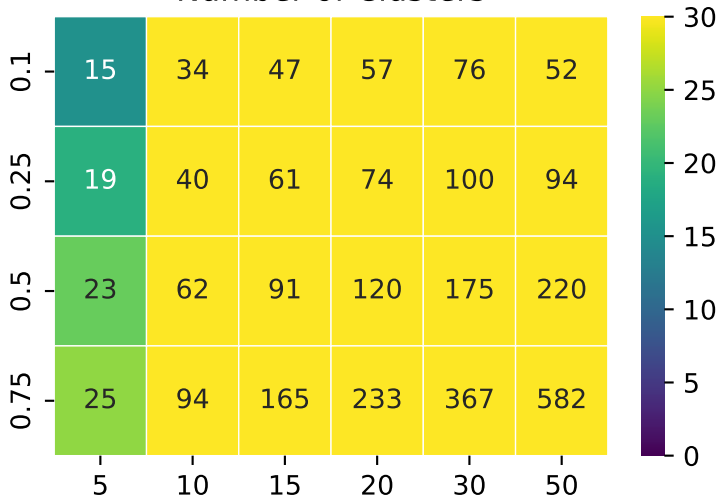
Projection: PCA (n_components=2)

Clustering: DBSCAN (eps=10.0)

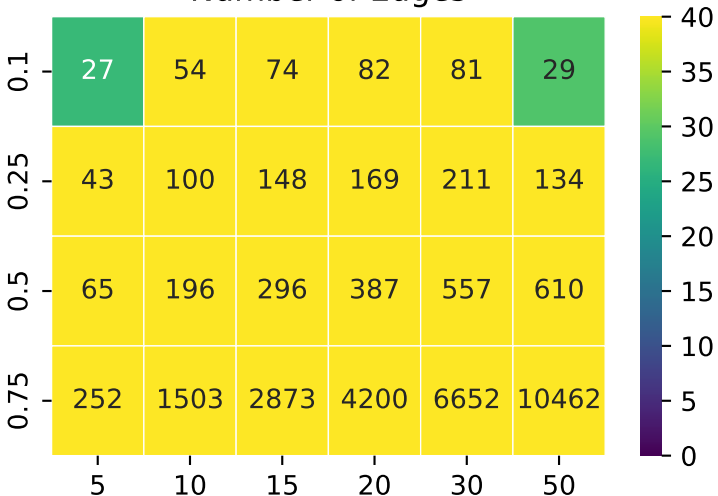
Unclustered samples



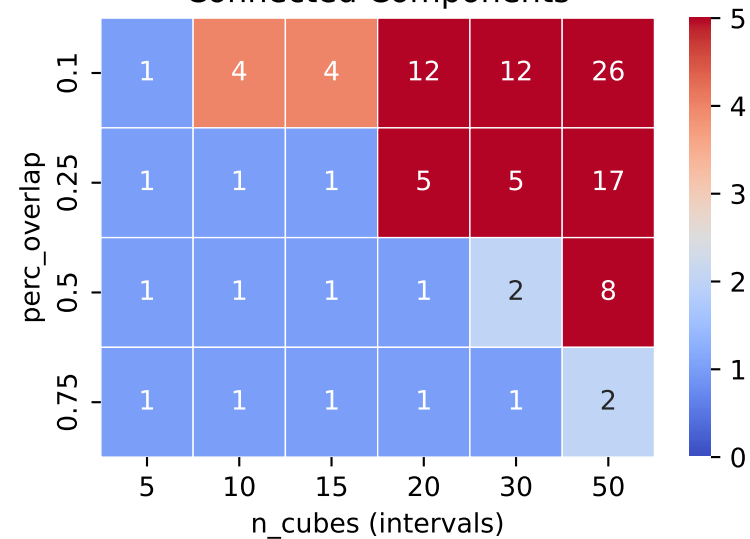
Number of Clusters



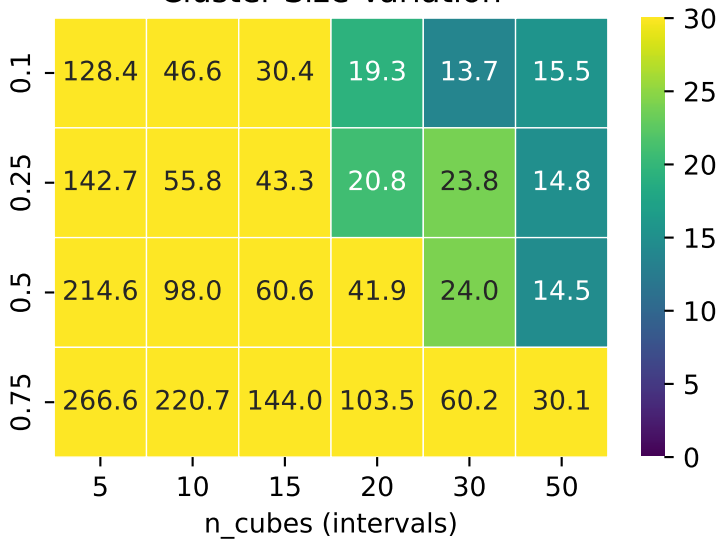
Number of Edges



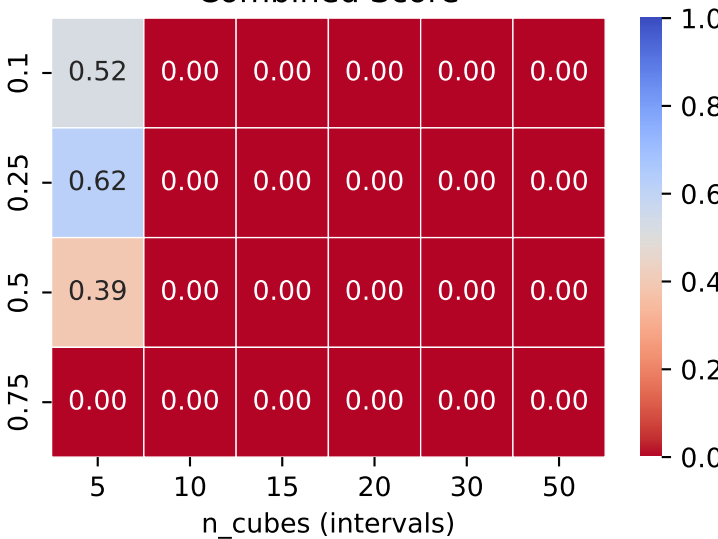
Connected Components



Cluster Size Variation

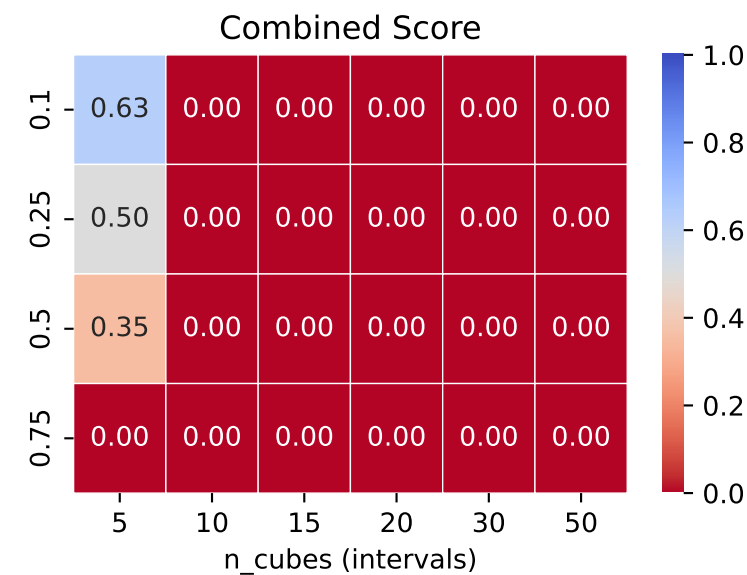
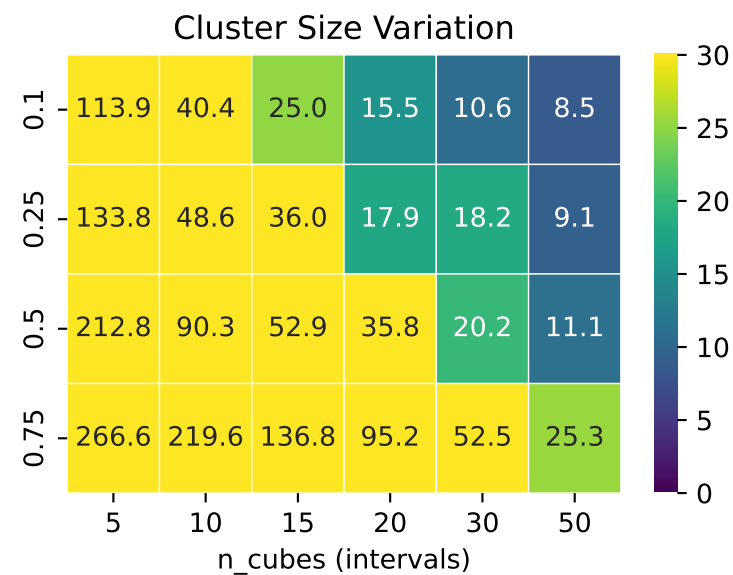
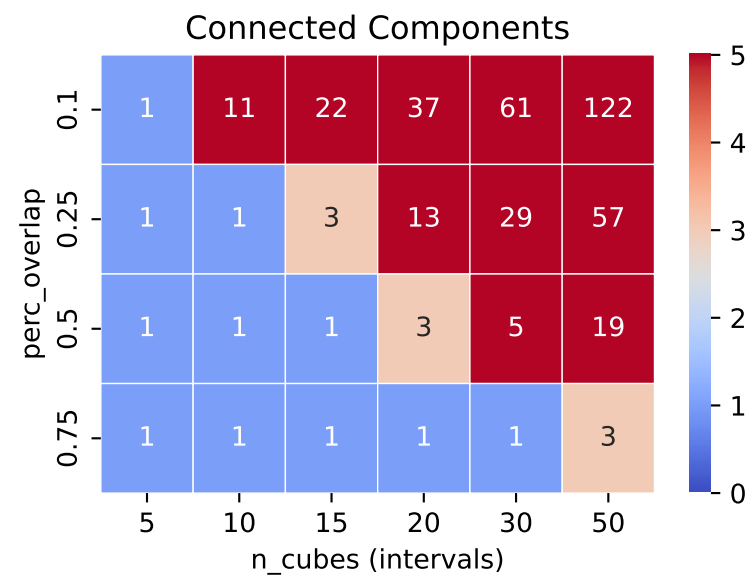
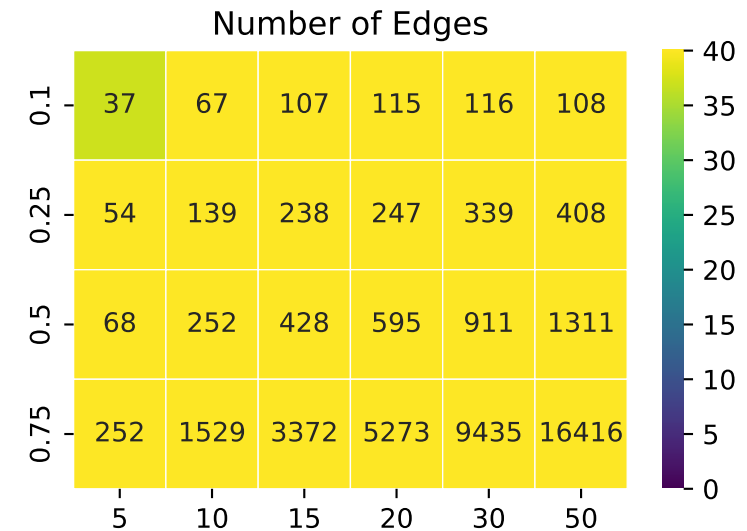
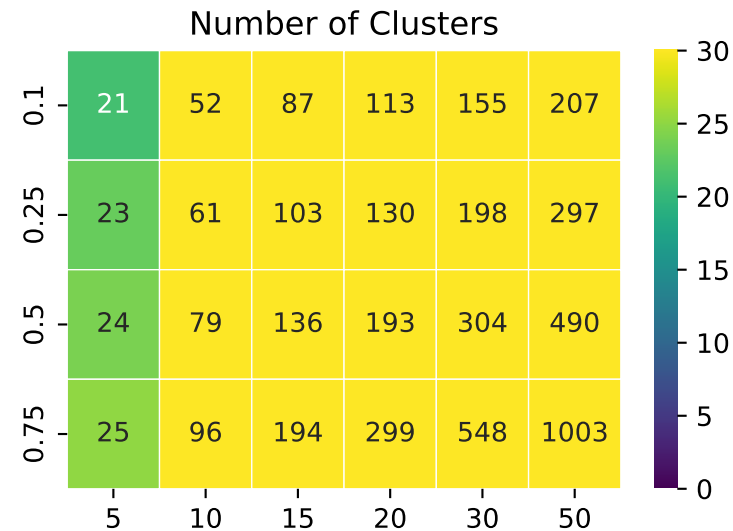
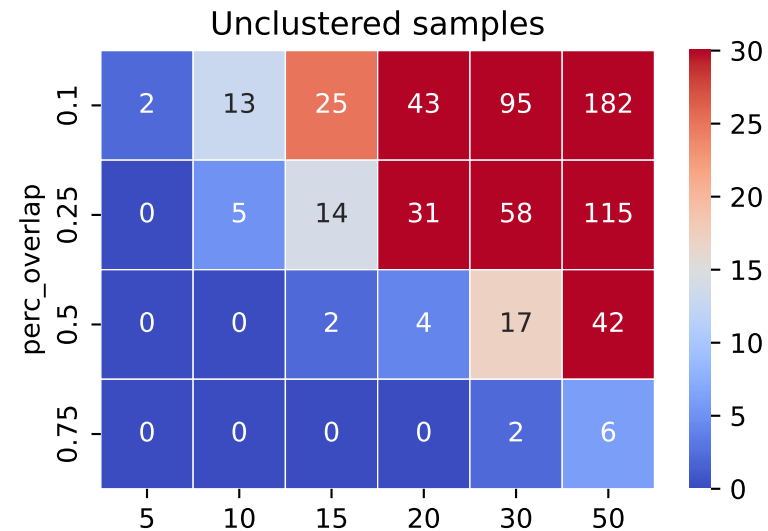


Combined Score



Projection: PCA (n_components=2)

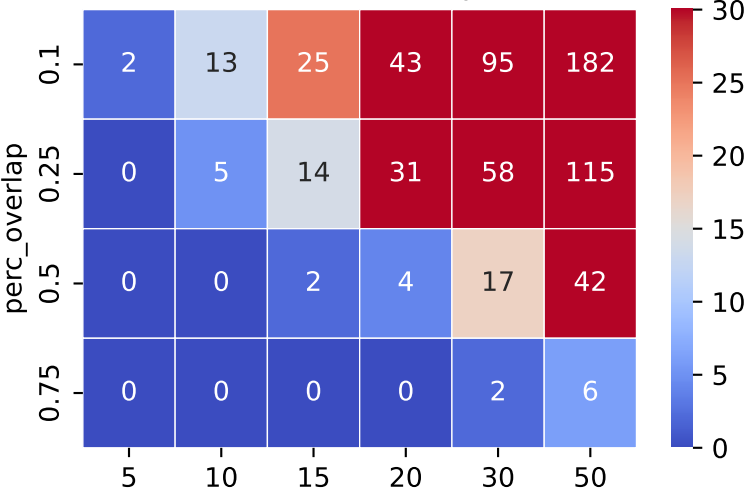
Clustering: Bayesian Gaussian Mixture (n_components=1)



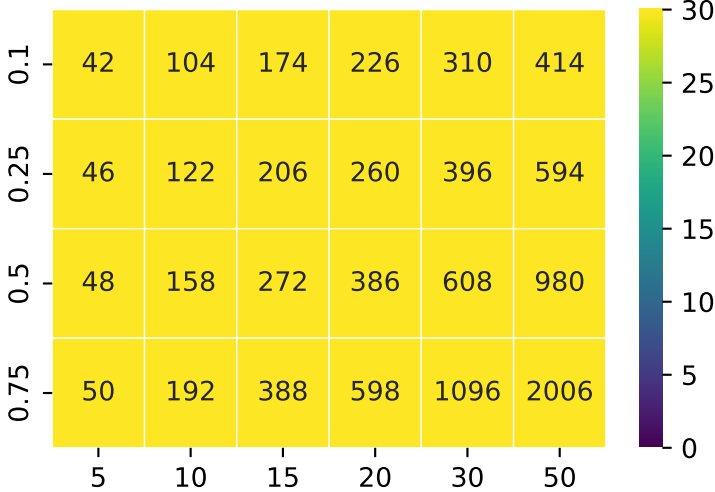
Projection: PCA (n_components=2)

Clustering: Bayesian Gaussian Mixture (n_components=2)

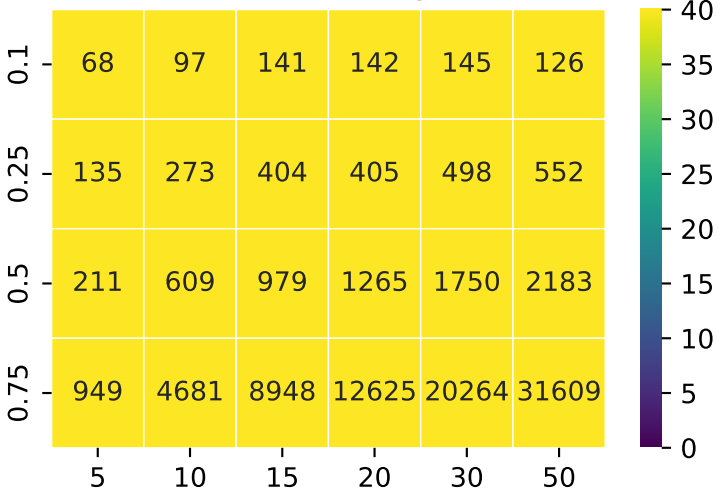
Unclustered samples



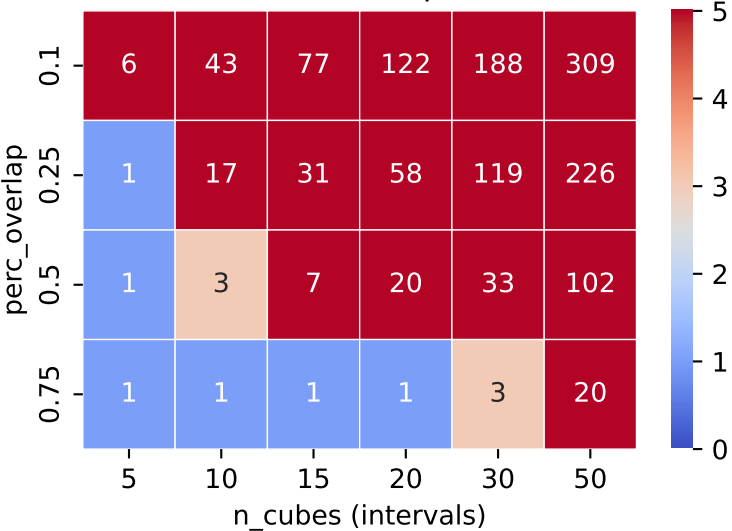
Number of Clusters



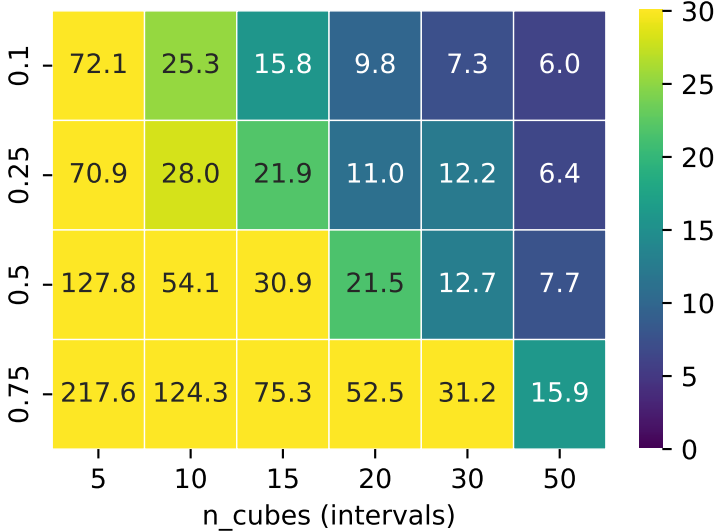
Number of Edges



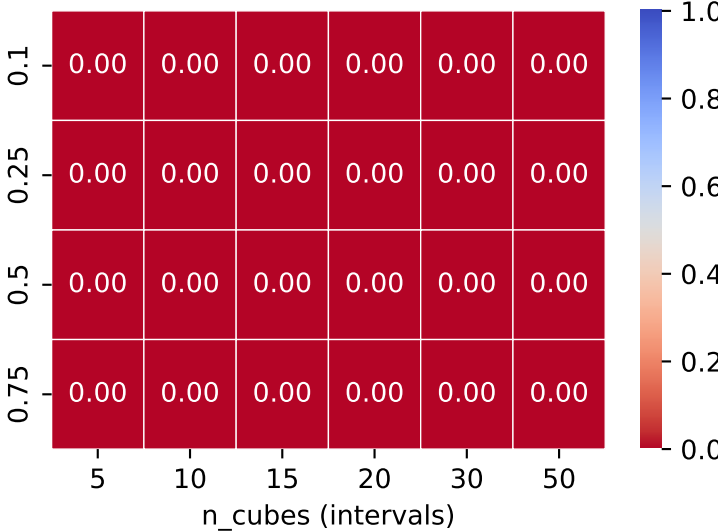
Connected Components



Cluster Size Variation



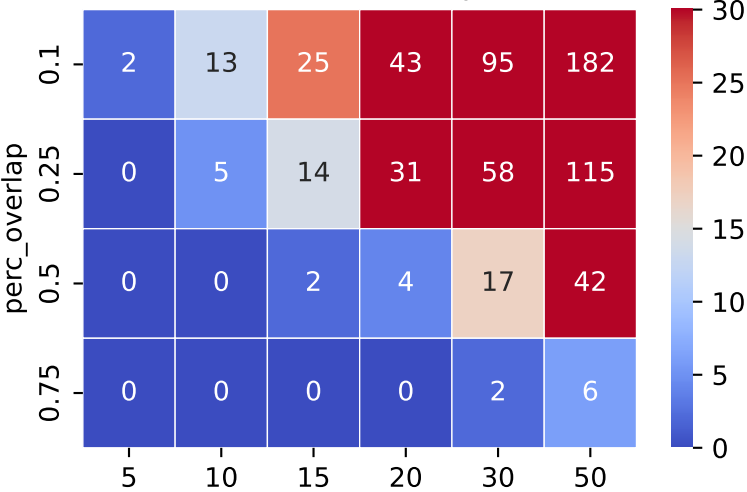
Combined Score



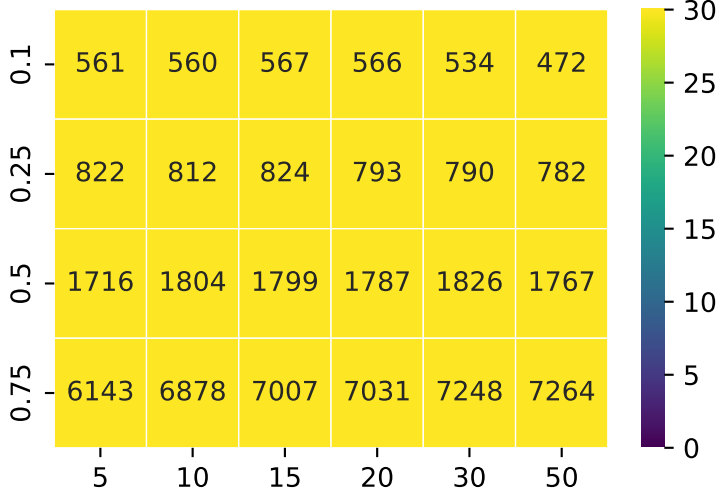
Projection: PCA (n_components=2)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=1.0)

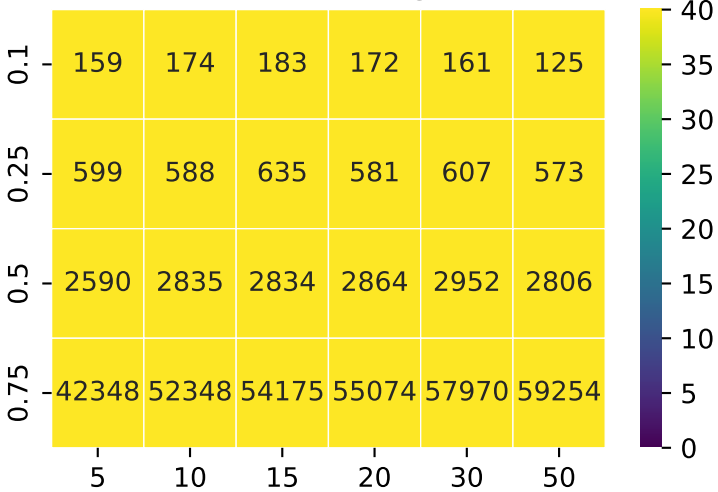
Unclustered samples



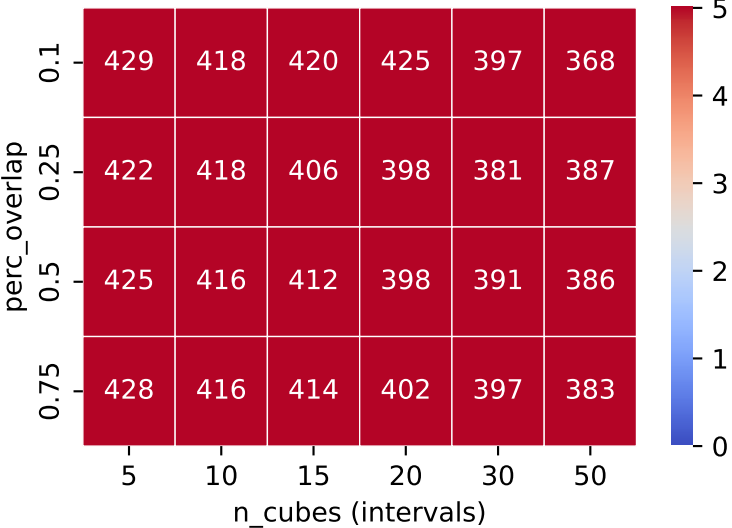
Number of Clusters



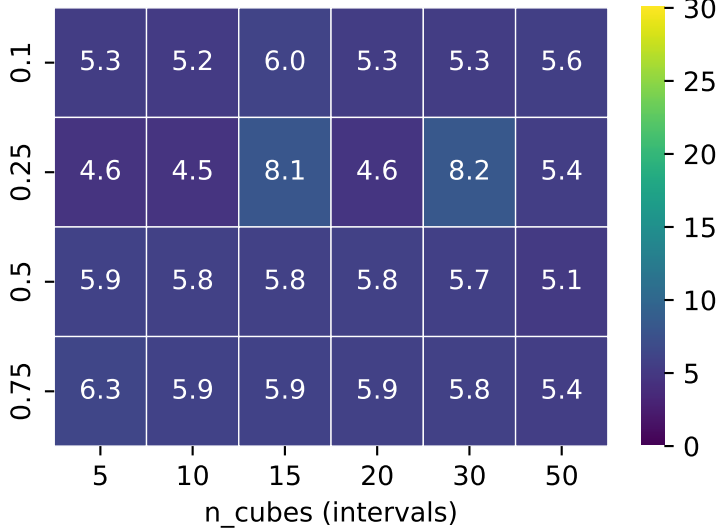
Number of Edges



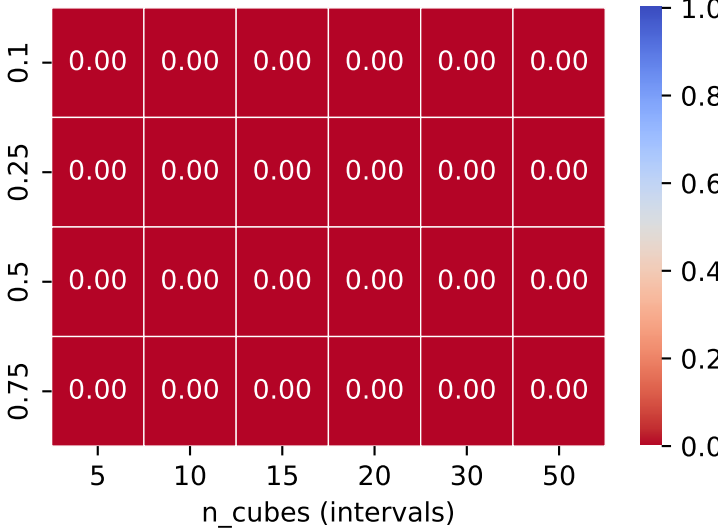
Connected Components



Cluster Size Variation



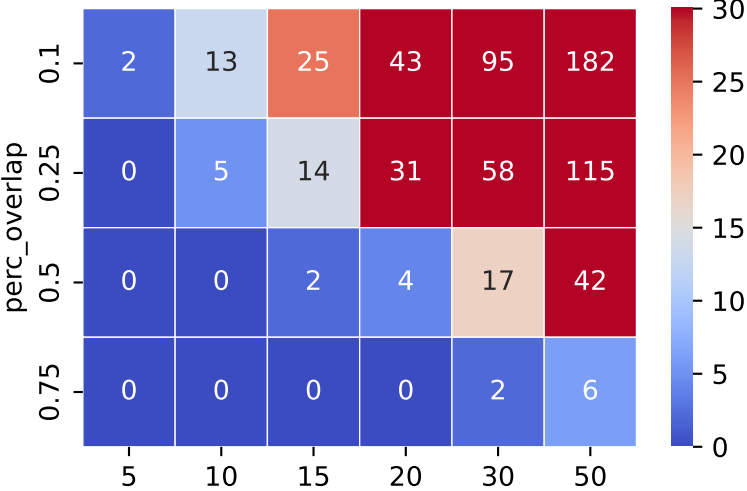
Combined Score



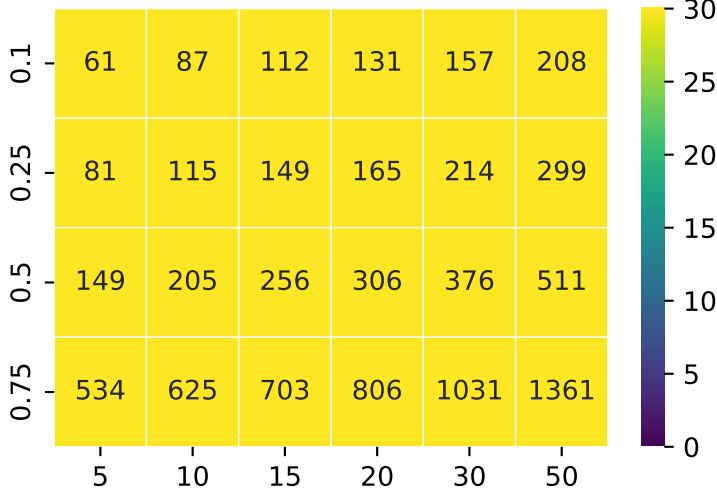
Projection: PCA (n_components=2)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=10.0)

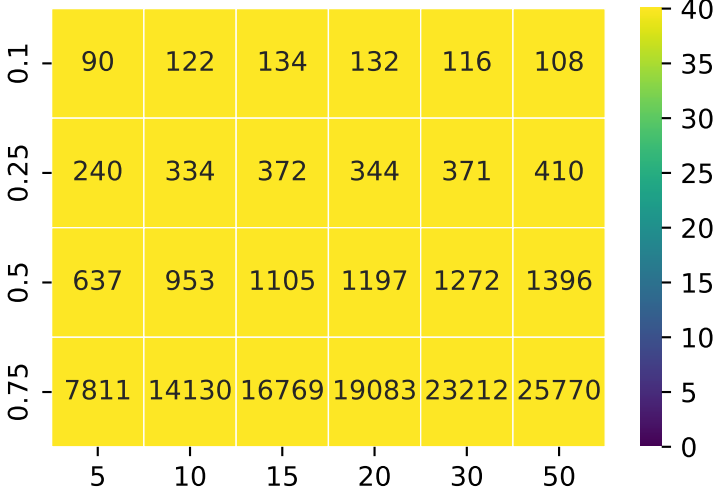
Unclustered samples



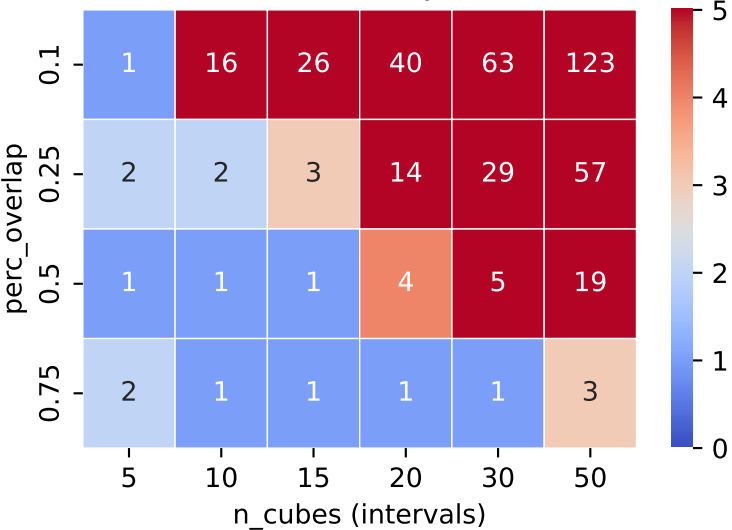
Number of Clusters



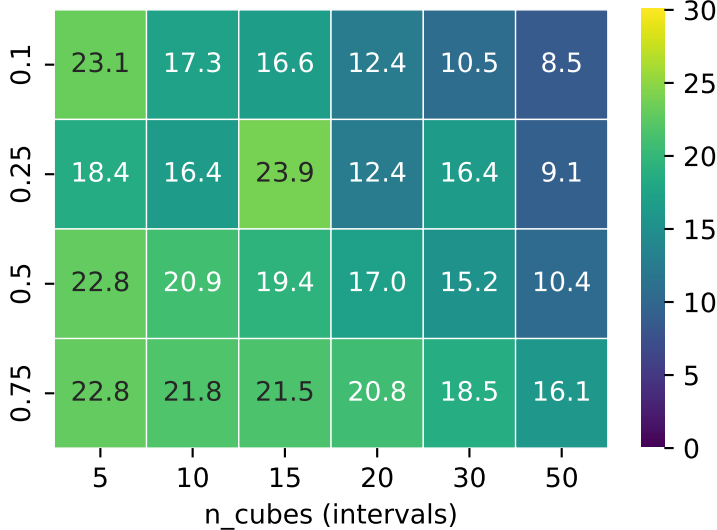
Number of Edges



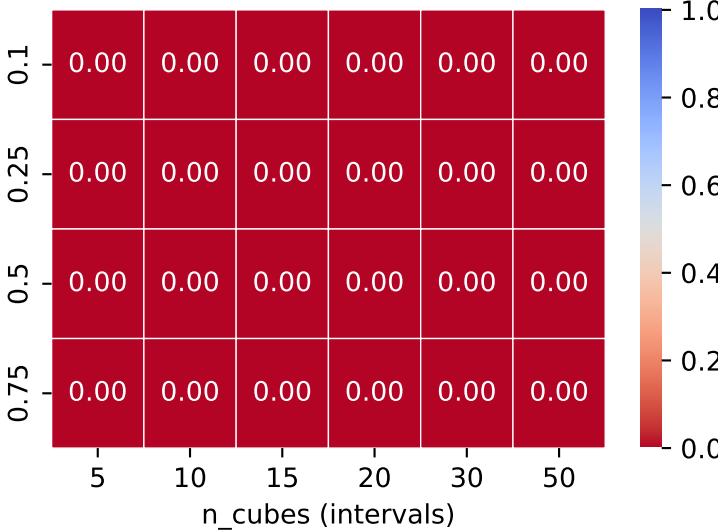
Connected Components



Cluster Size Variation

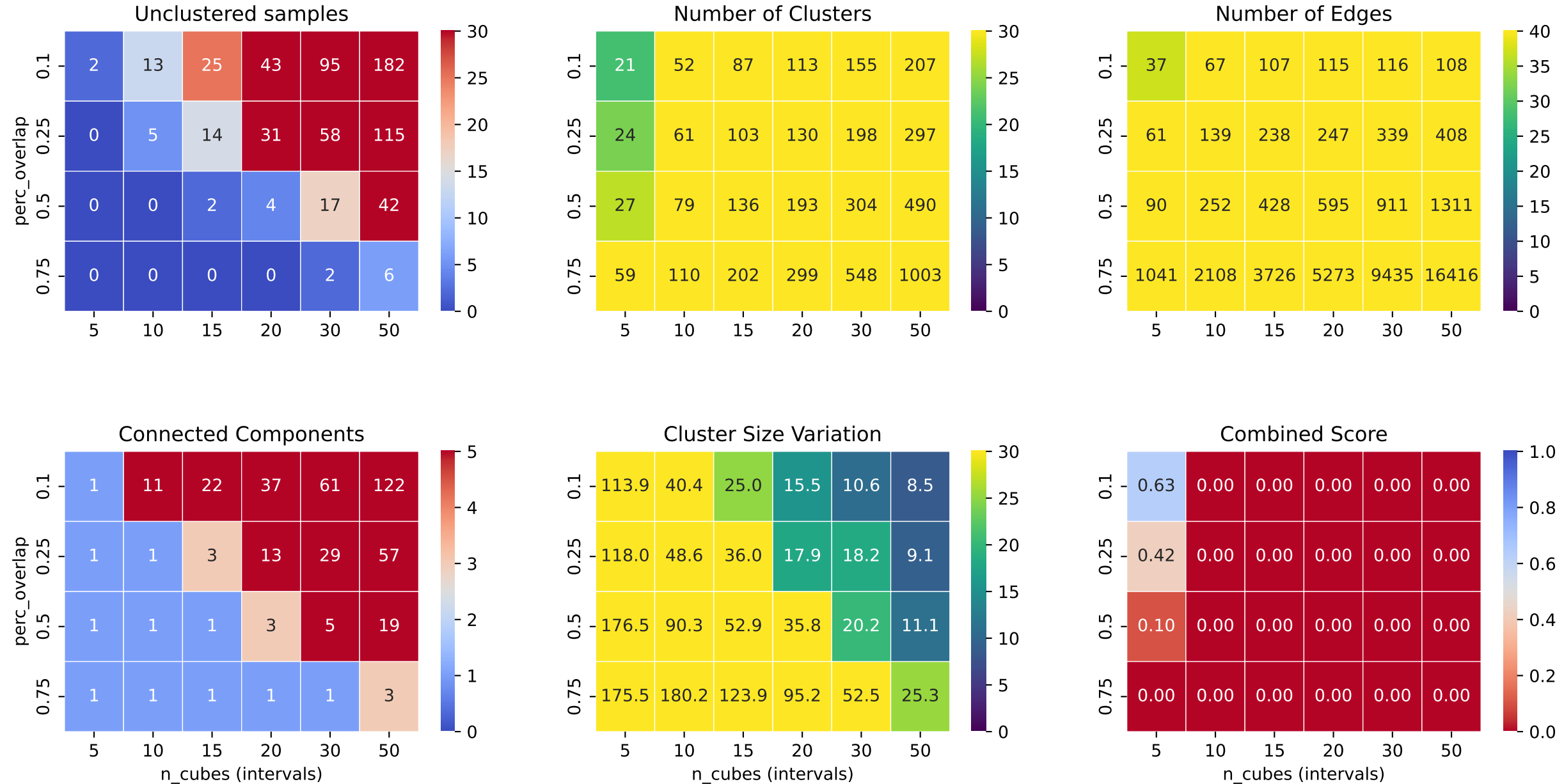


Combined Score



Projection: PCA (n_components=2)

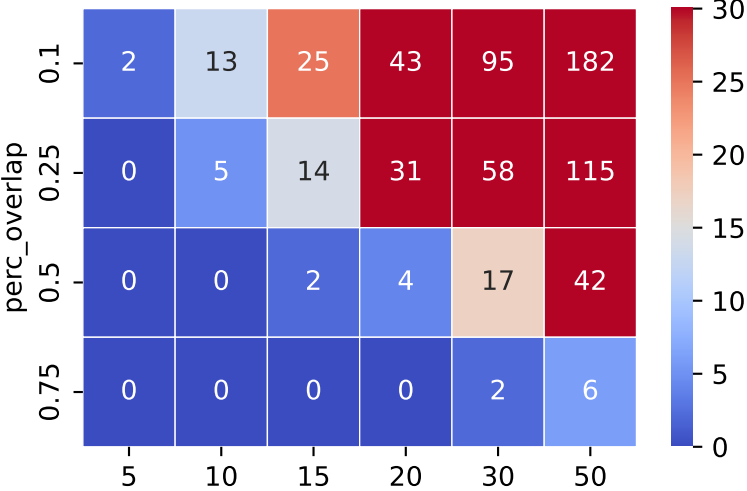
Clustering: Hierarchical Agglomerative Clustering (distance_threshold=30.0)



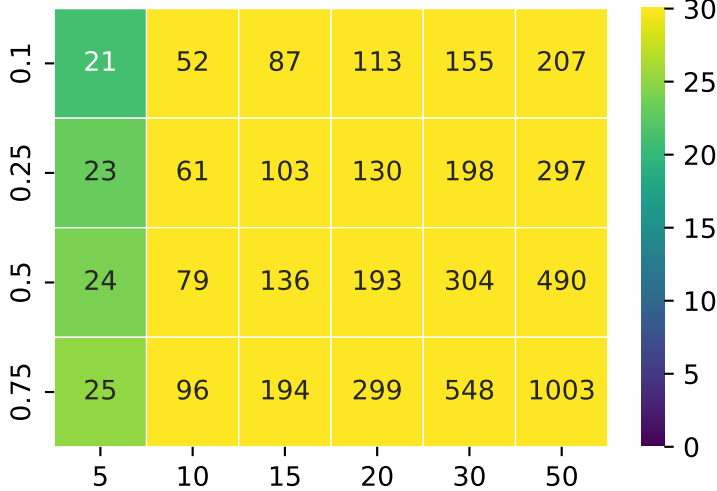
Projection: PCA (n_components=2)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

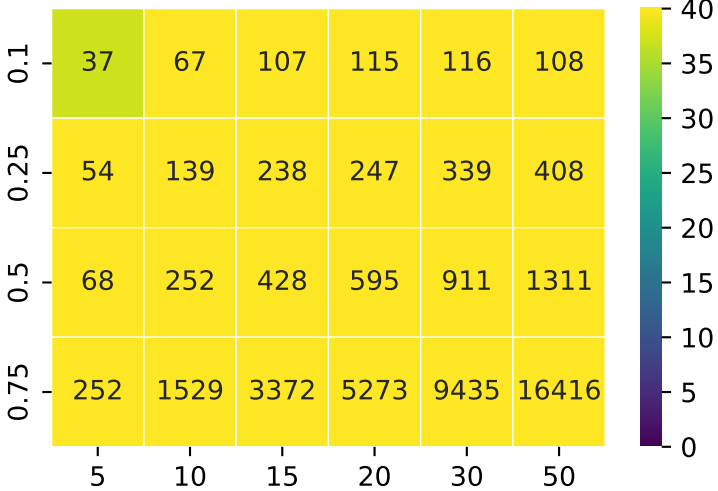
Unclustered samples



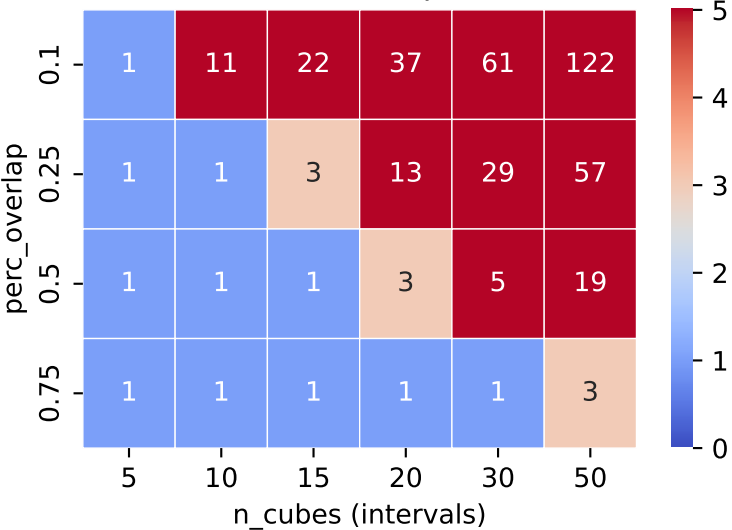
Number of Clusters



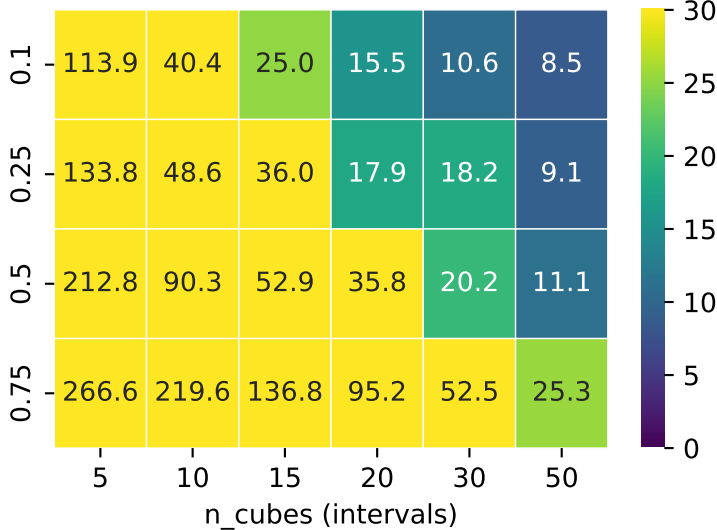
Number of Edges



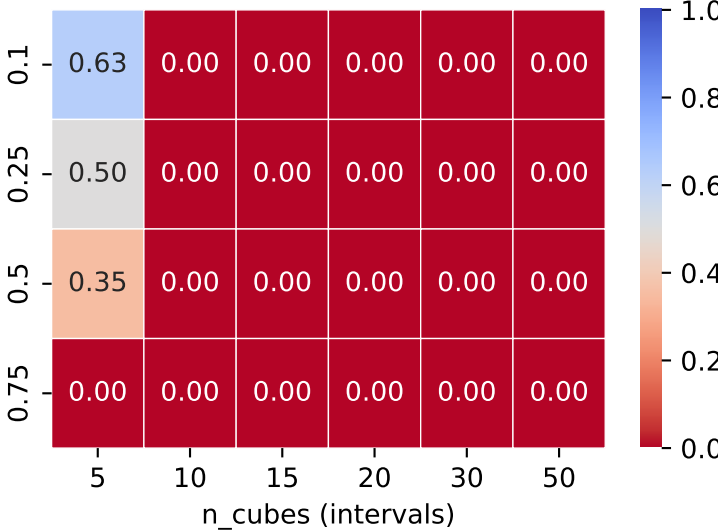
Connected Components



Cluster Size Variation



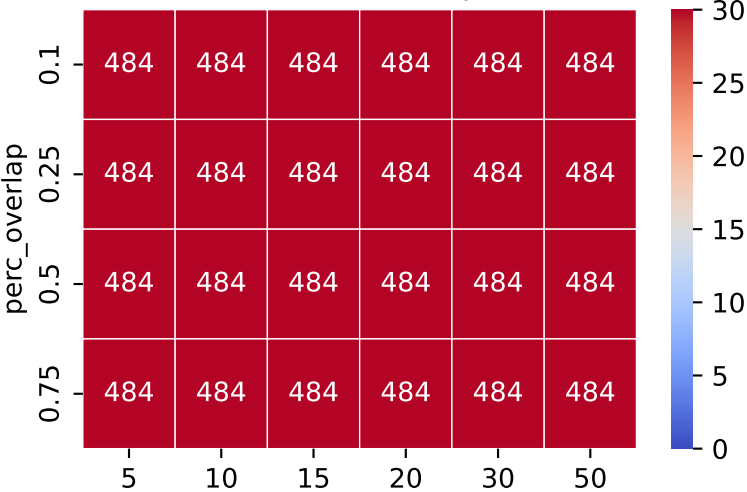
Combined Score



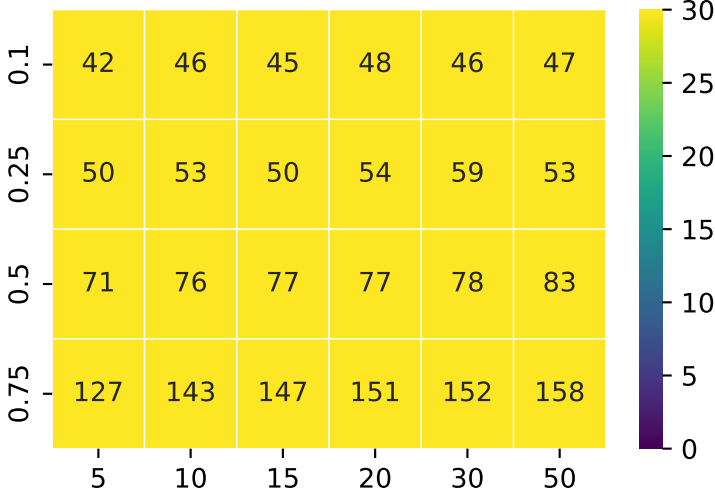
Projection: t-SNE (n_components=1)

Clustering: DBSCAN (eps=1.0)

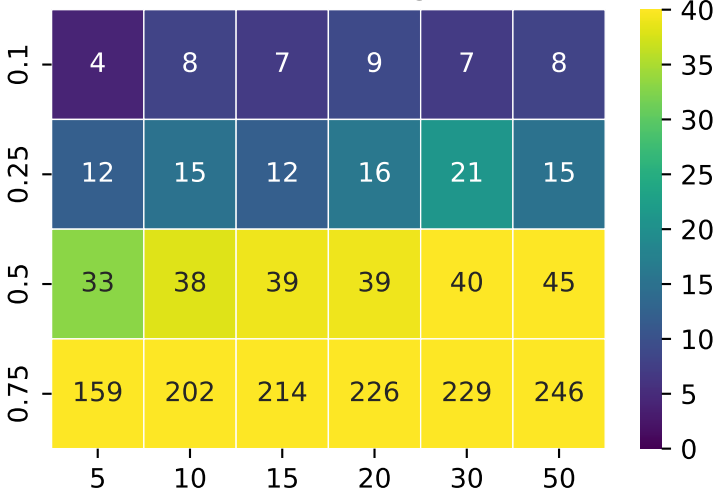
Unclustered samples



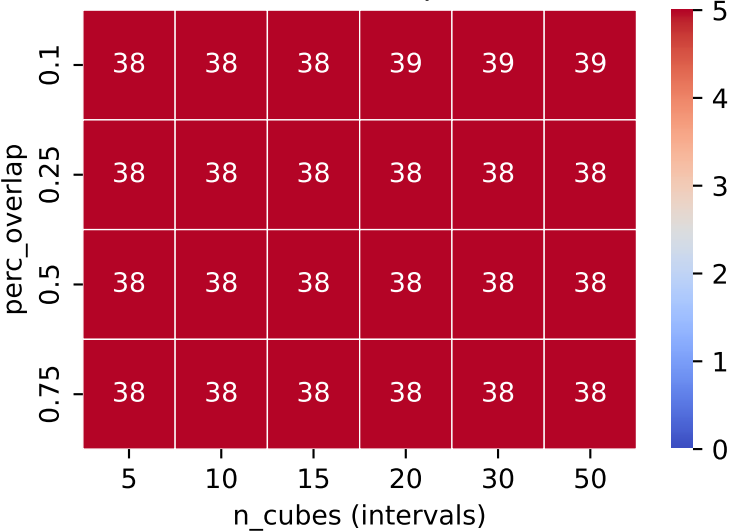
Number of Clusters



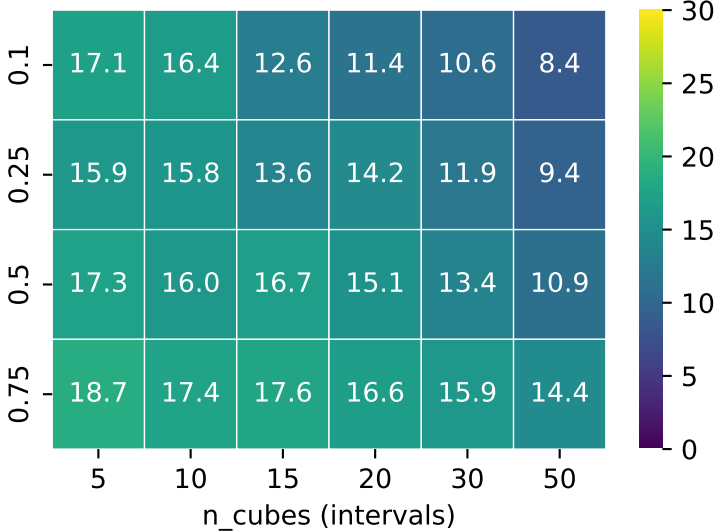
Number of Edges



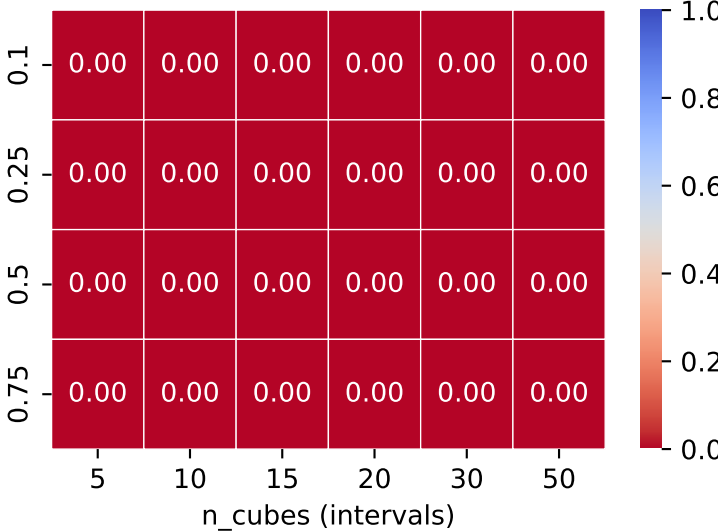
Connected Components



Cluster Size Variation



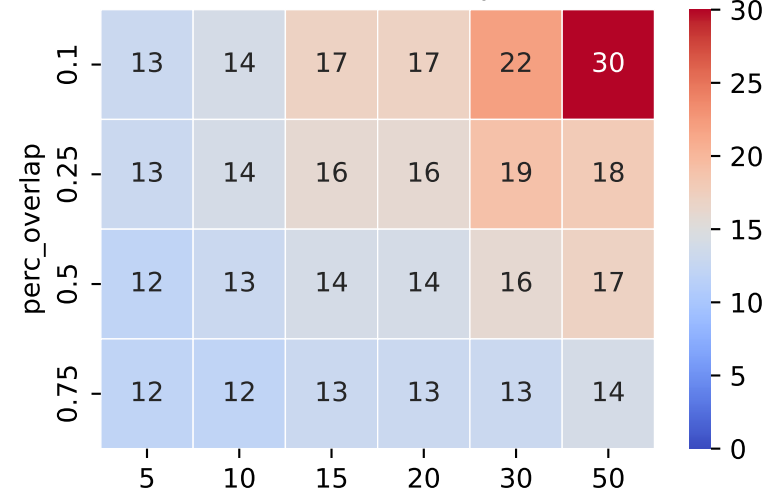
Combined Score



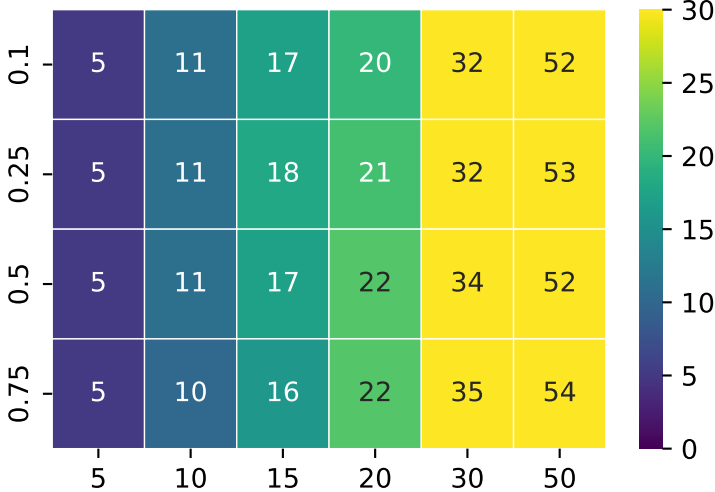
Projection: t-SNE (n_components=1)

Clustering: DBSCAN (eps=3.0)

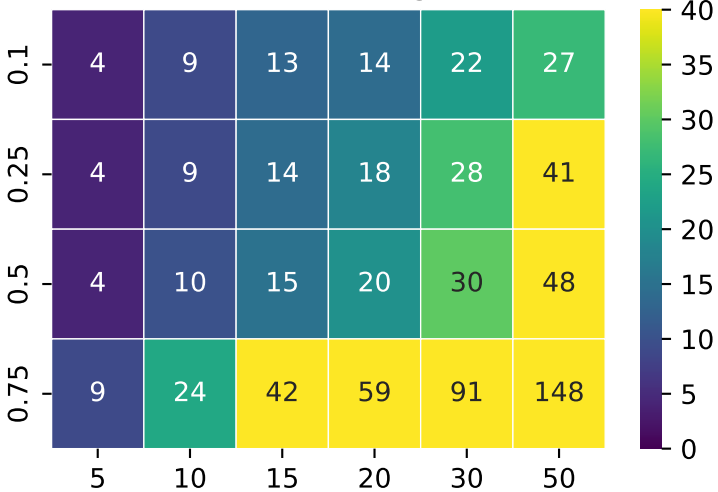
Unclustered samples



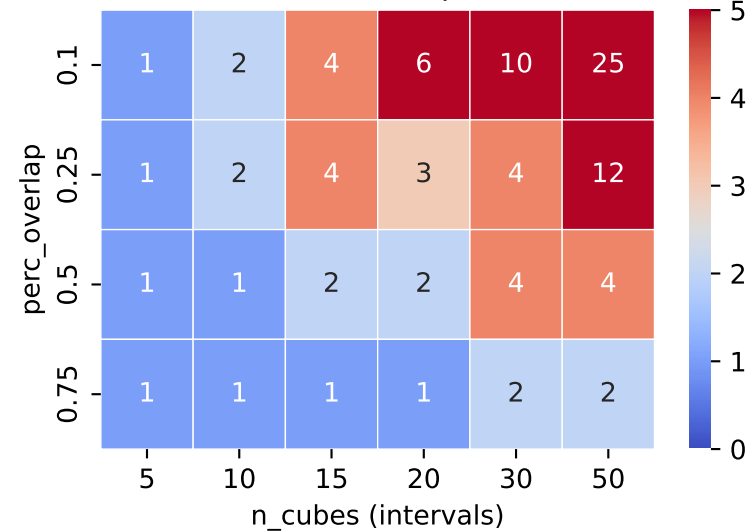
Number of Clusters



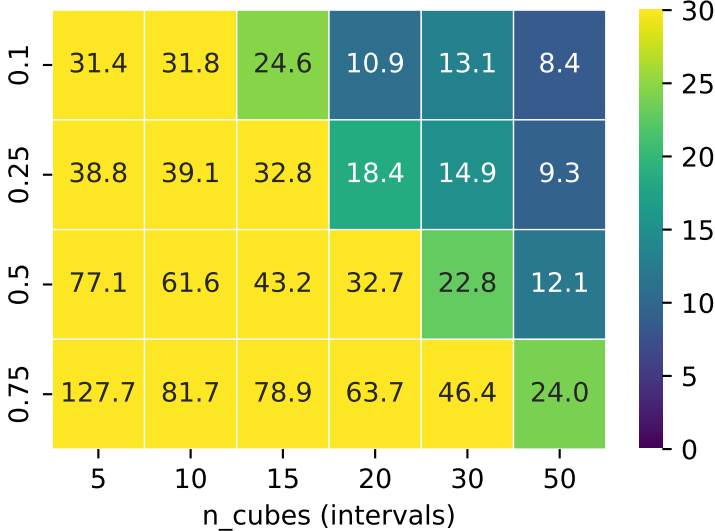
Number of Edges



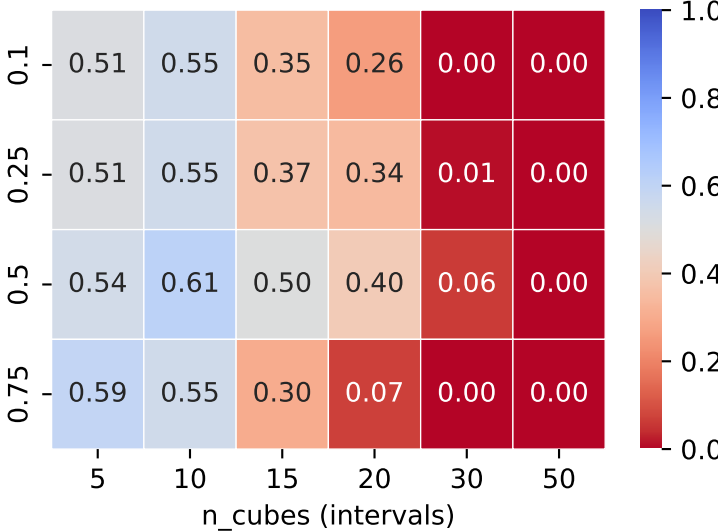
Connected Components



Cluster Size Variation



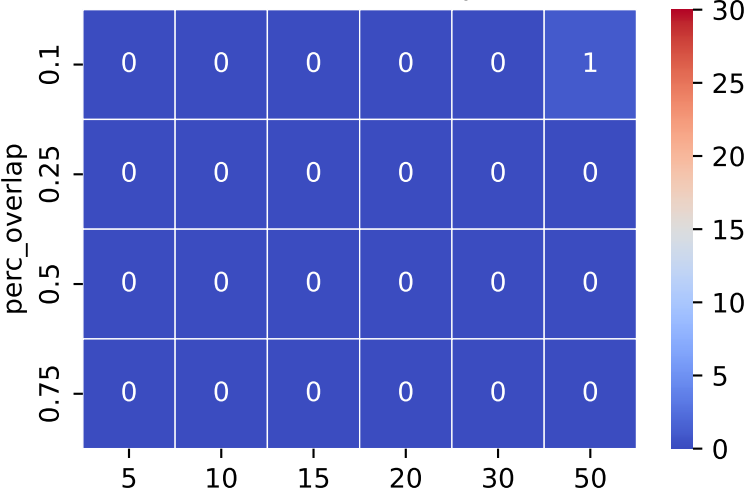
Combined Score



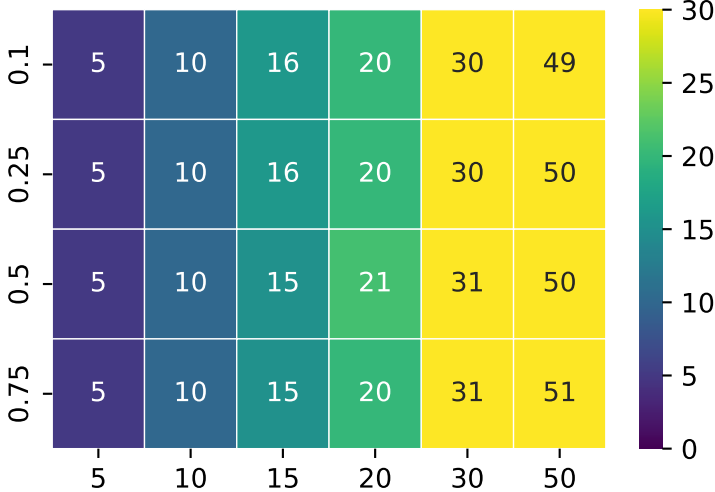
Projection: t-SNE (n_components=1)

Clustering: DBSCAN (eps=5.0)

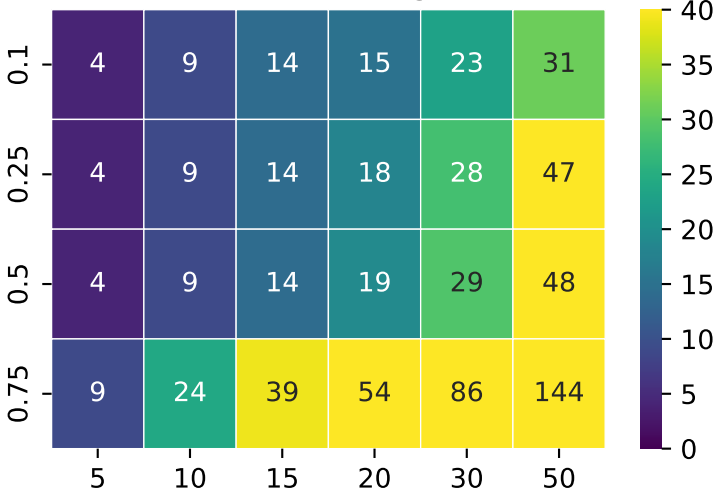
Unclustered samples



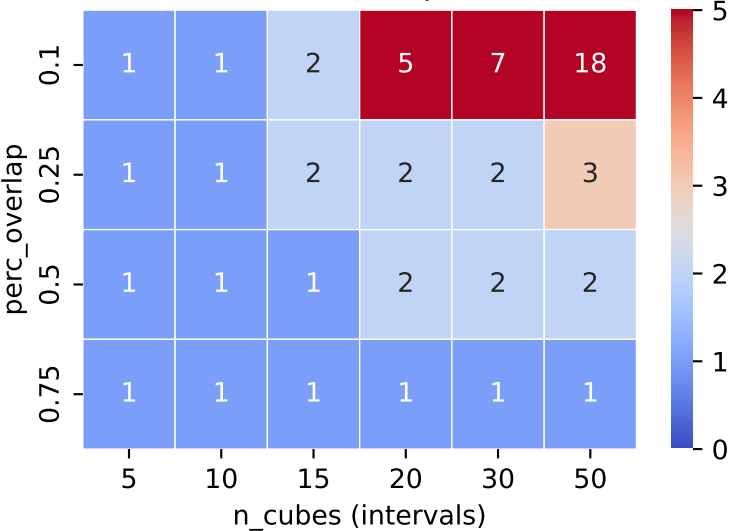
Number of Clusters



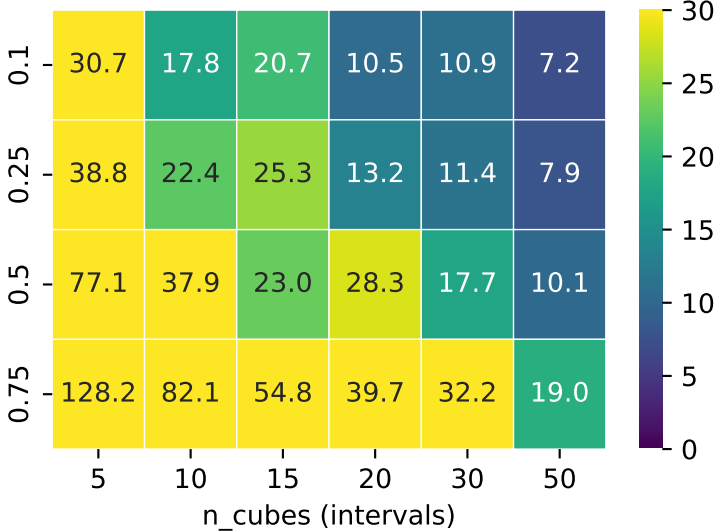
Number of Edges



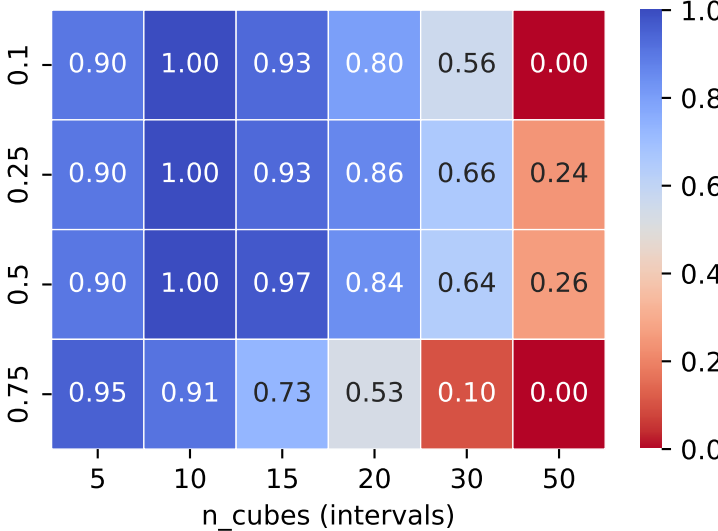
Connected Components



Cluster Size Variation



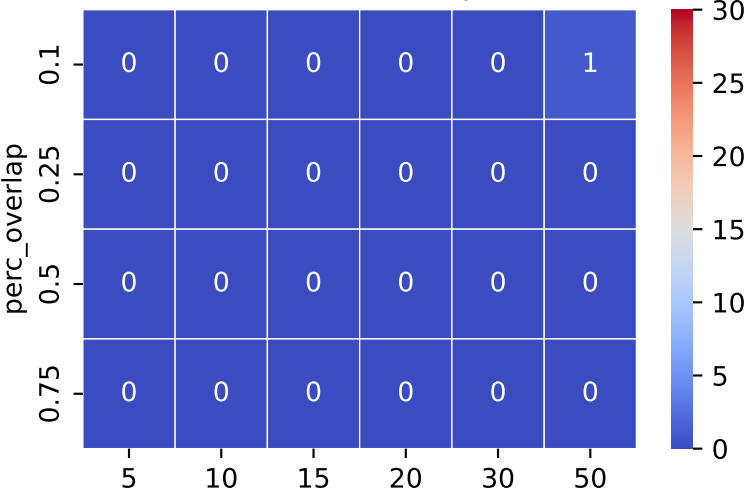
Combined Score



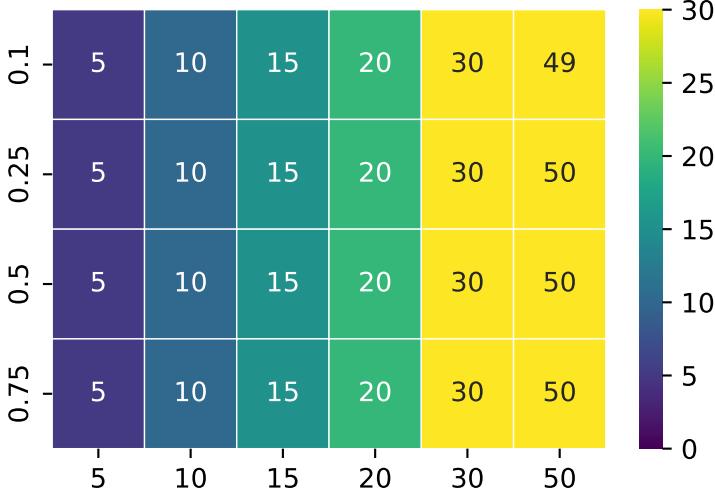
Projection: t-SNE (n_components=1)

Clustering: DBSCAN (eps=10.0)

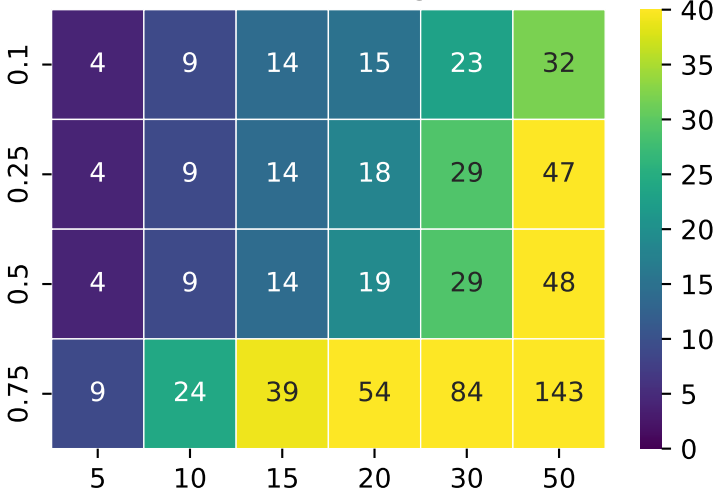
Unclustered samples



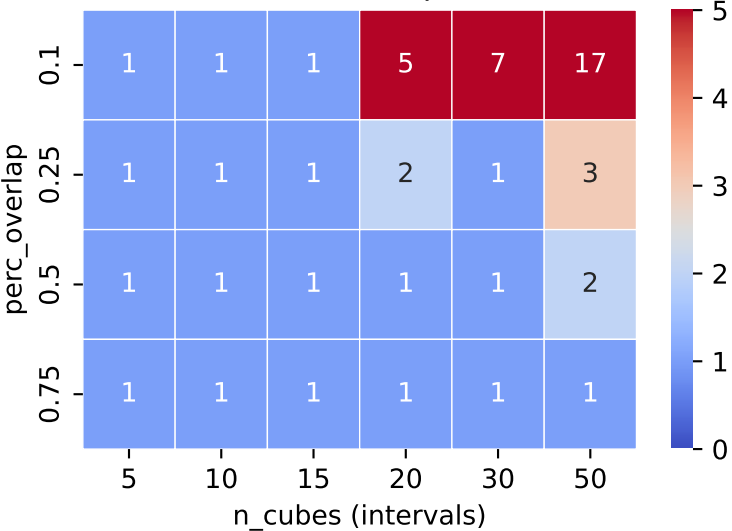
Number of Clusters



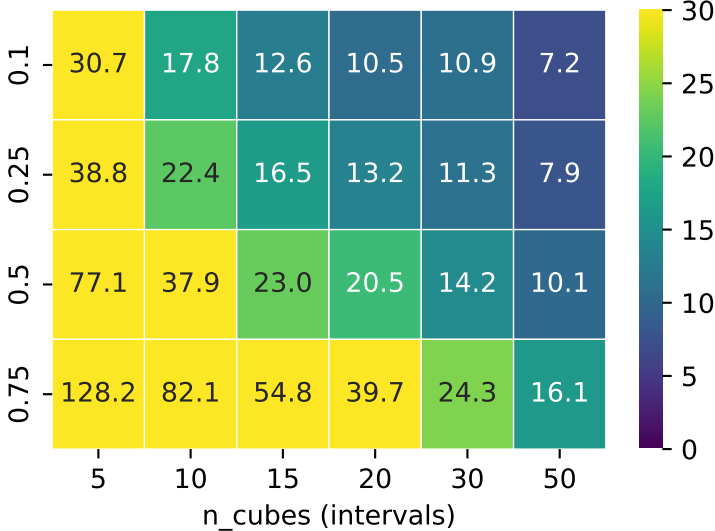
Number of Edges



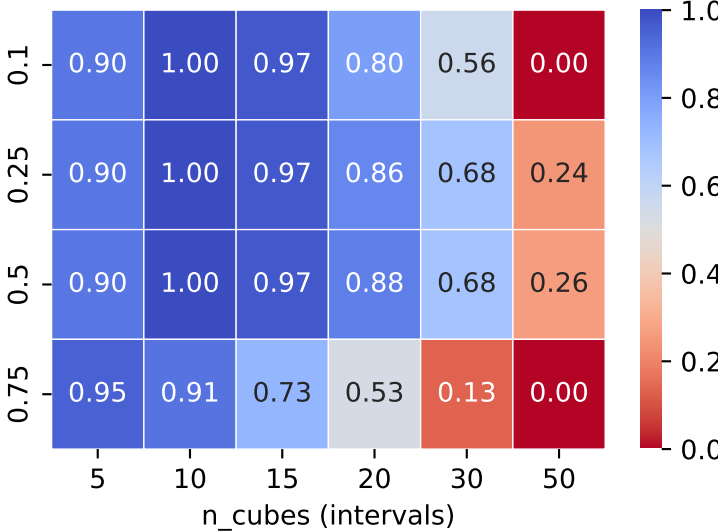
Connected Components



Cluster Size Variation

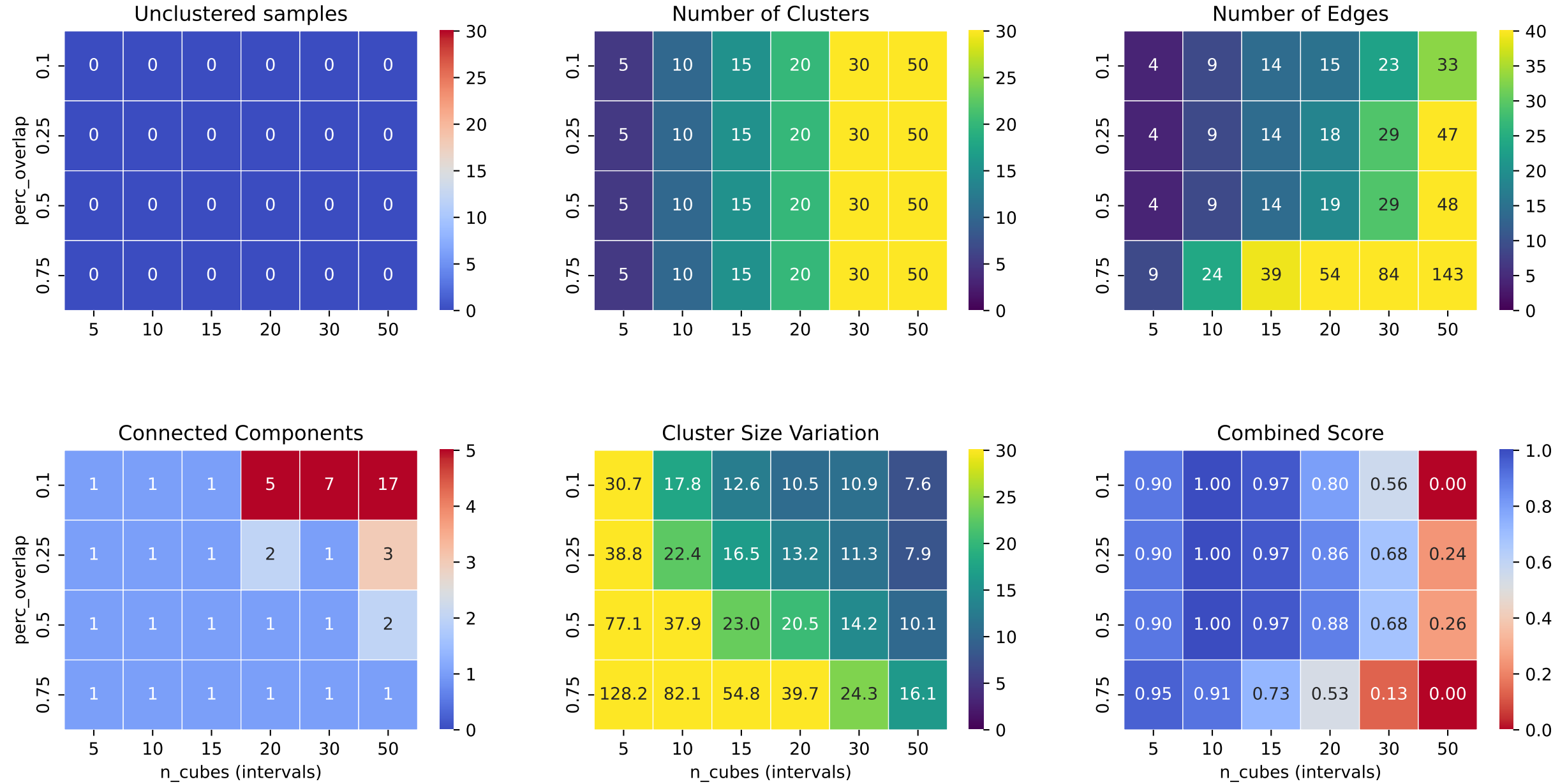


Combined Score



Projection: t-SNE (n_components=1)

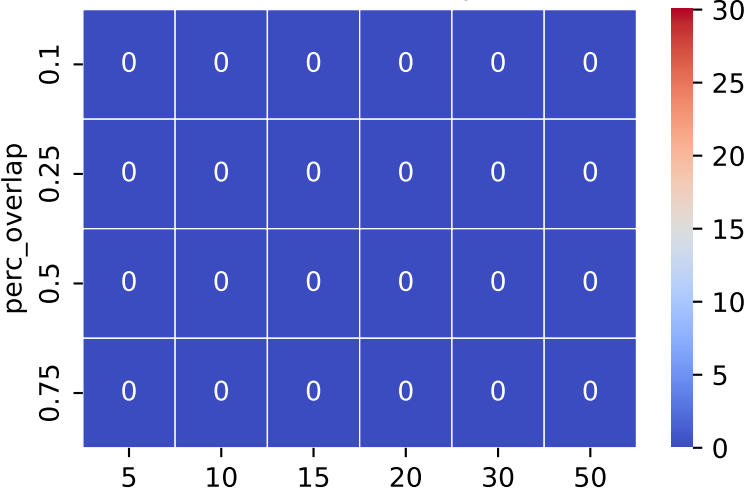
Clustering: Bayesian Gaussian Mixture (n_components=1)



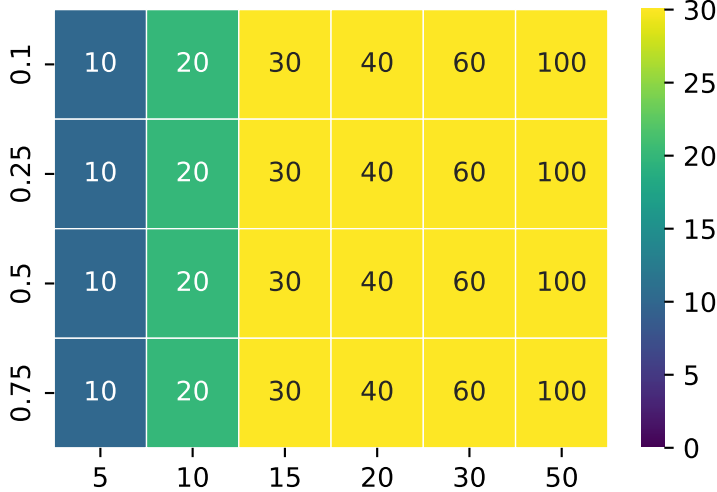
Projection: t-SNE (n_components=1)

Clustering: Bayesian Gaussian Mixture (n_components=2)

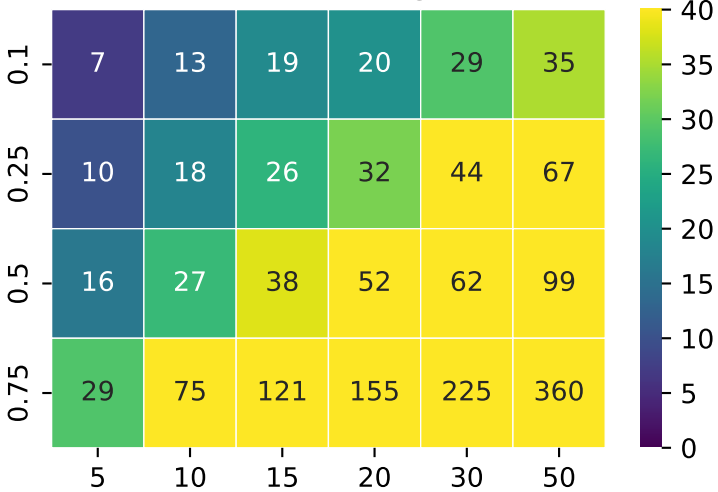
Unclustered samples



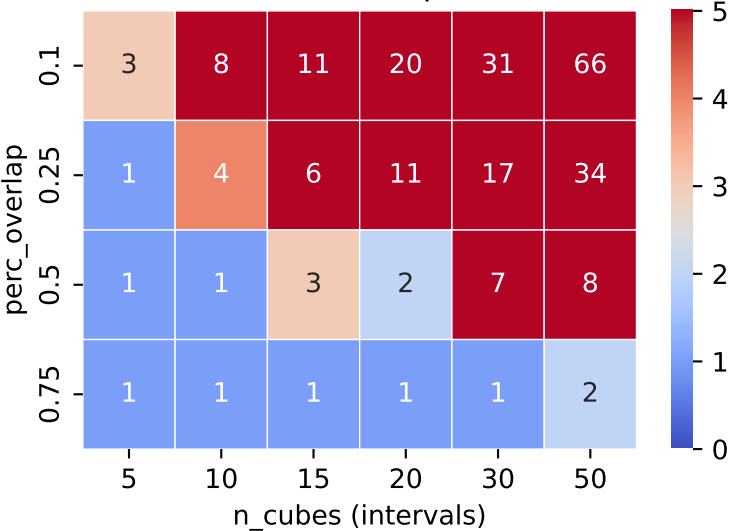
Number of Clusters



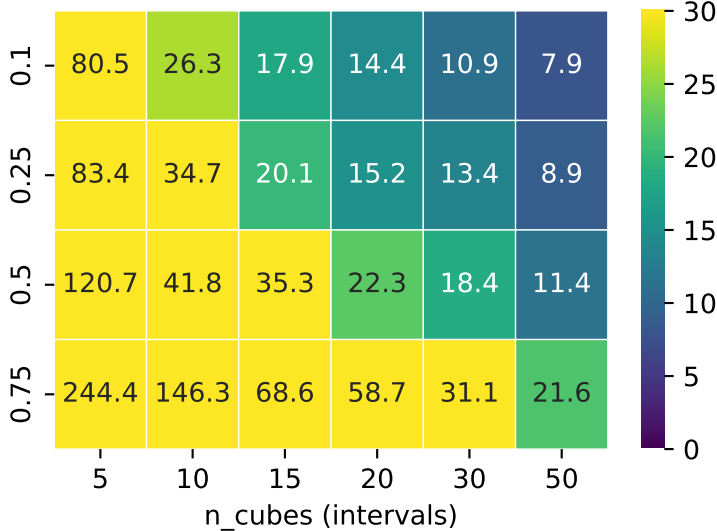
Number of Edges



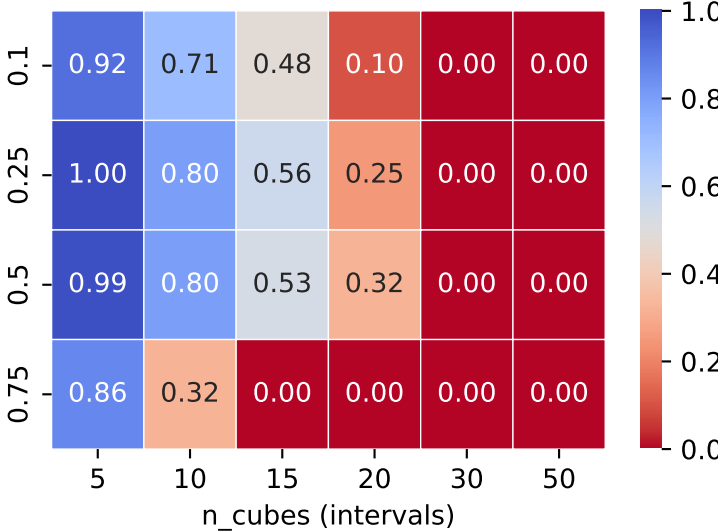
Connected Components



Cluster Size Variation



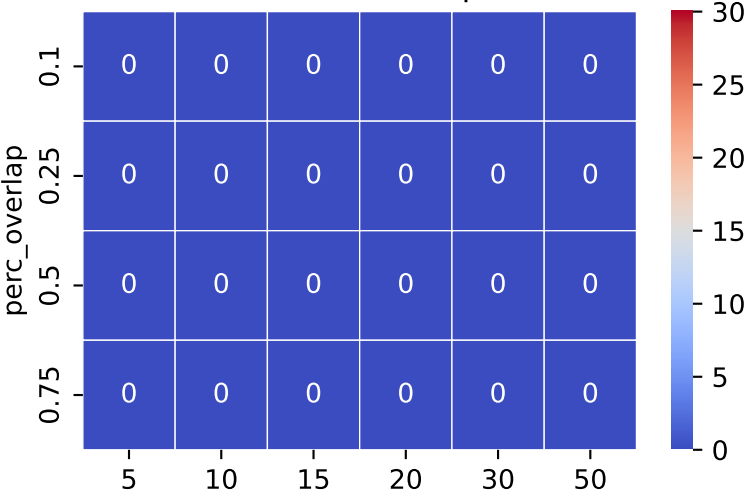
Combined Score



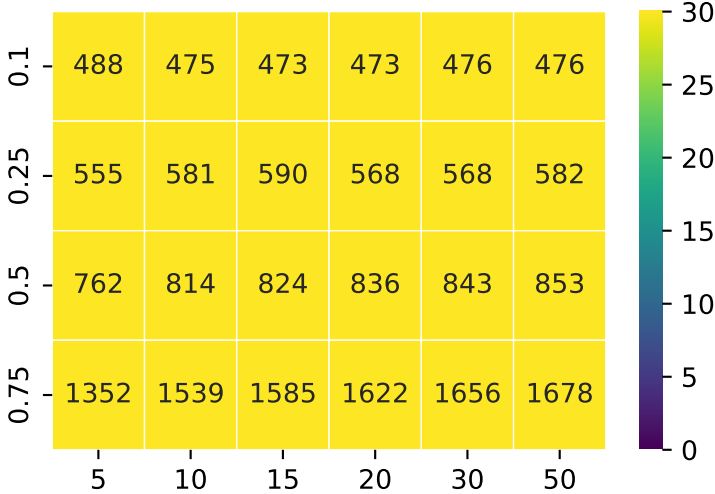
Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=1.0)

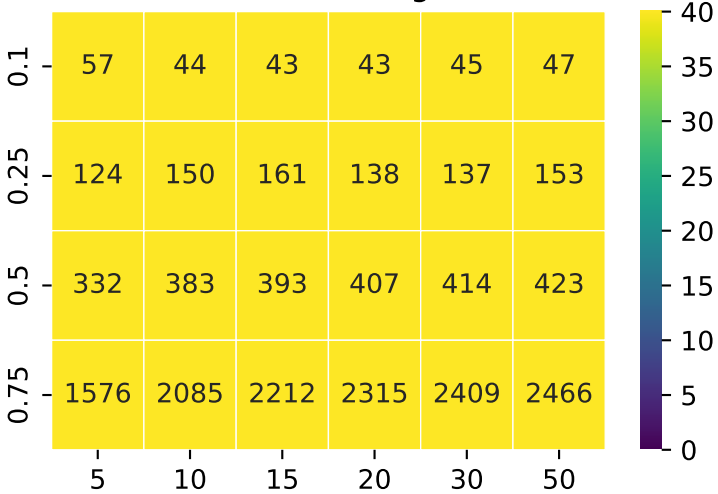
Unclustered samples



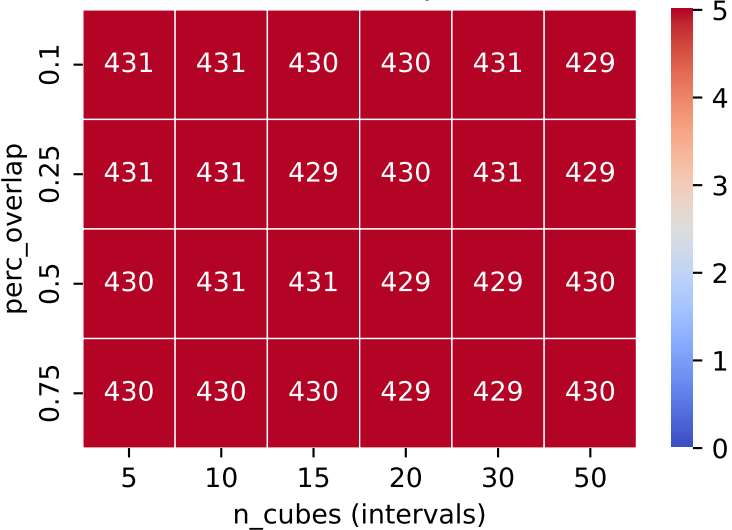
Number of Clusters



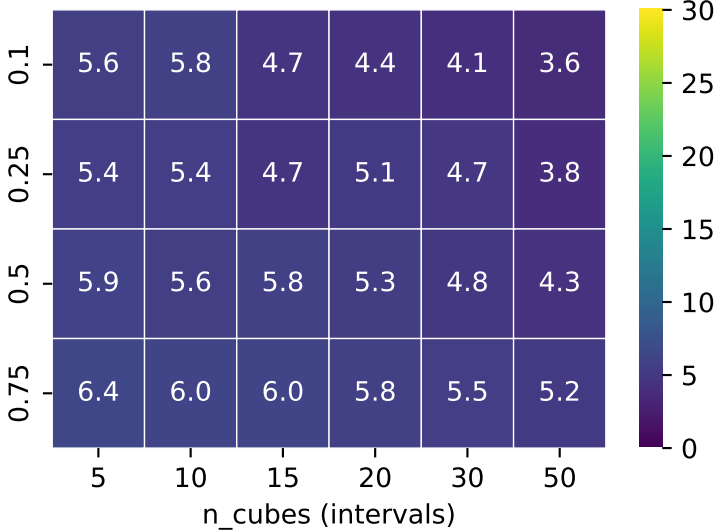
Number of Edges



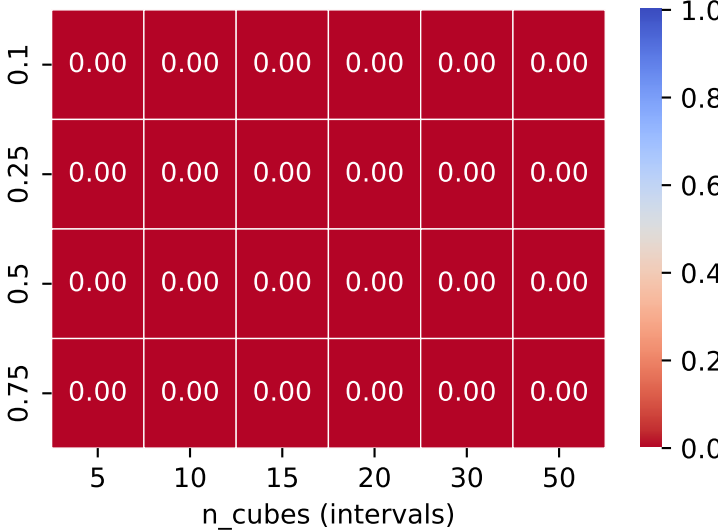
Connected Components



Cluster Size Variation

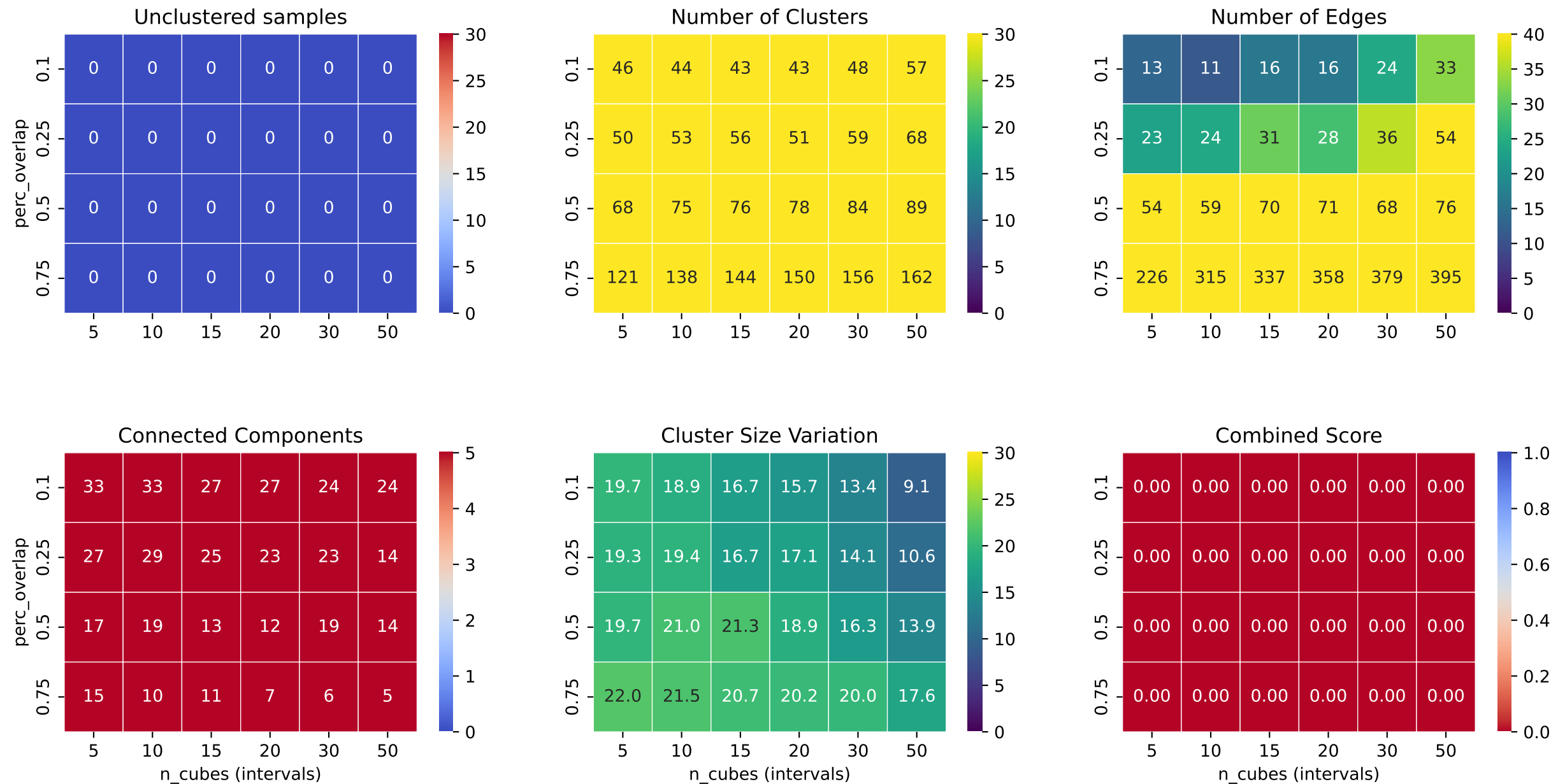


Combined Score



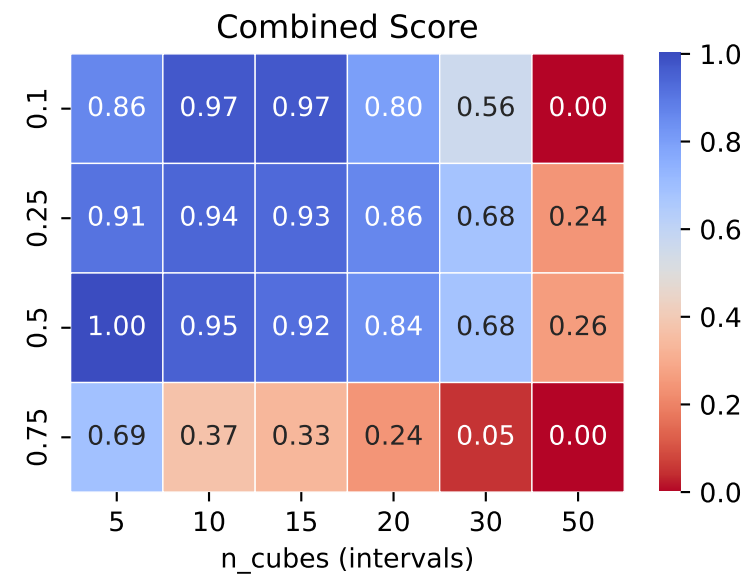
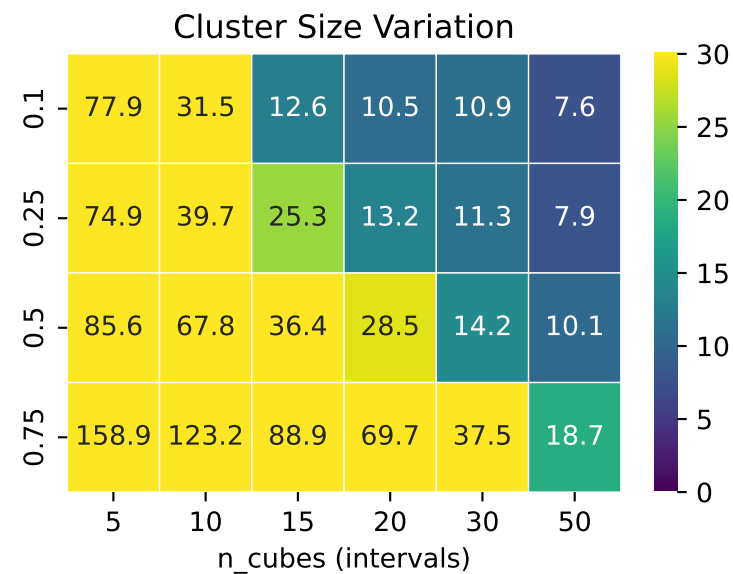
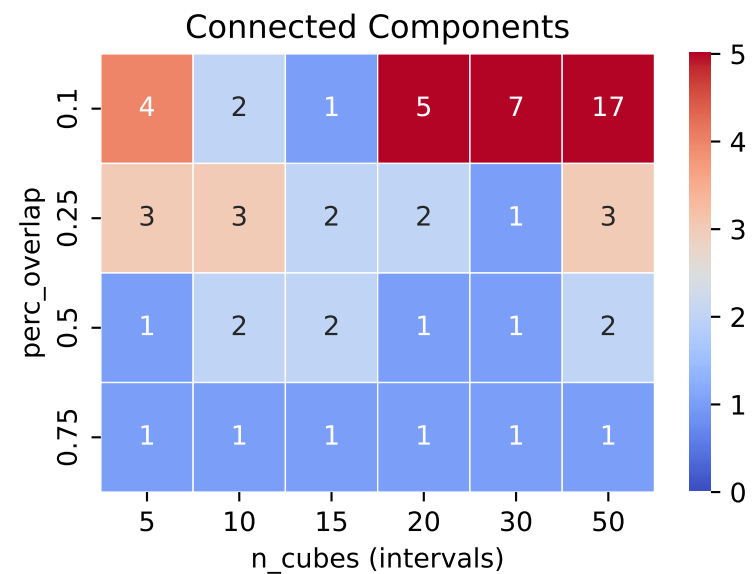
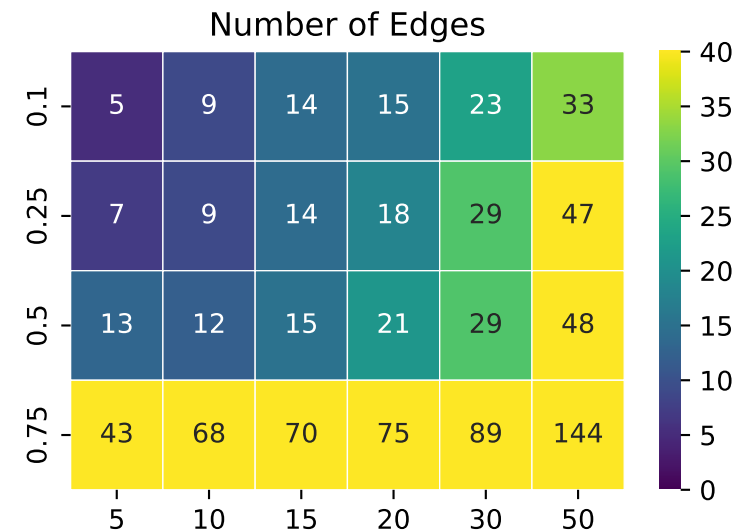
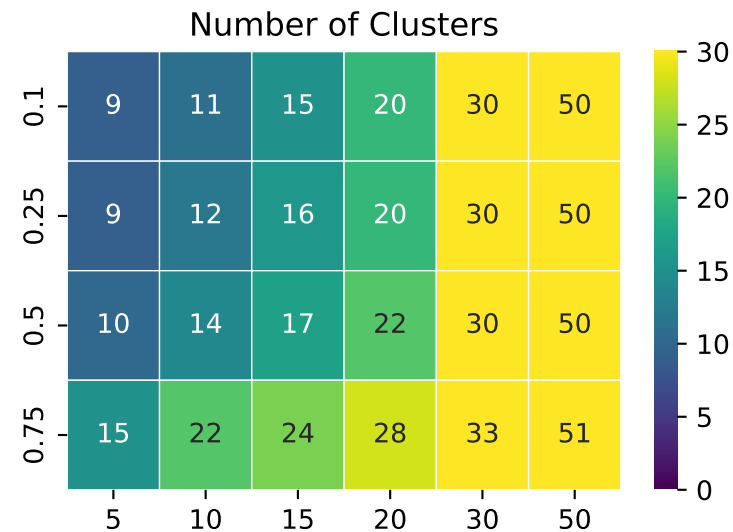
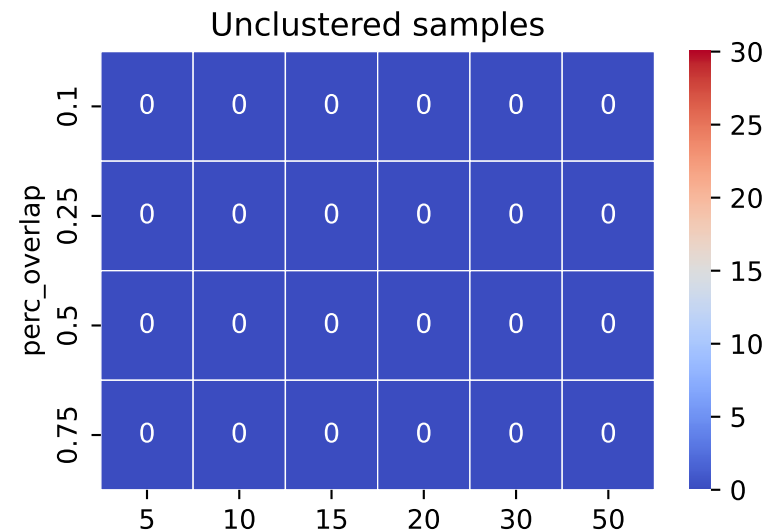
Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=10.0)



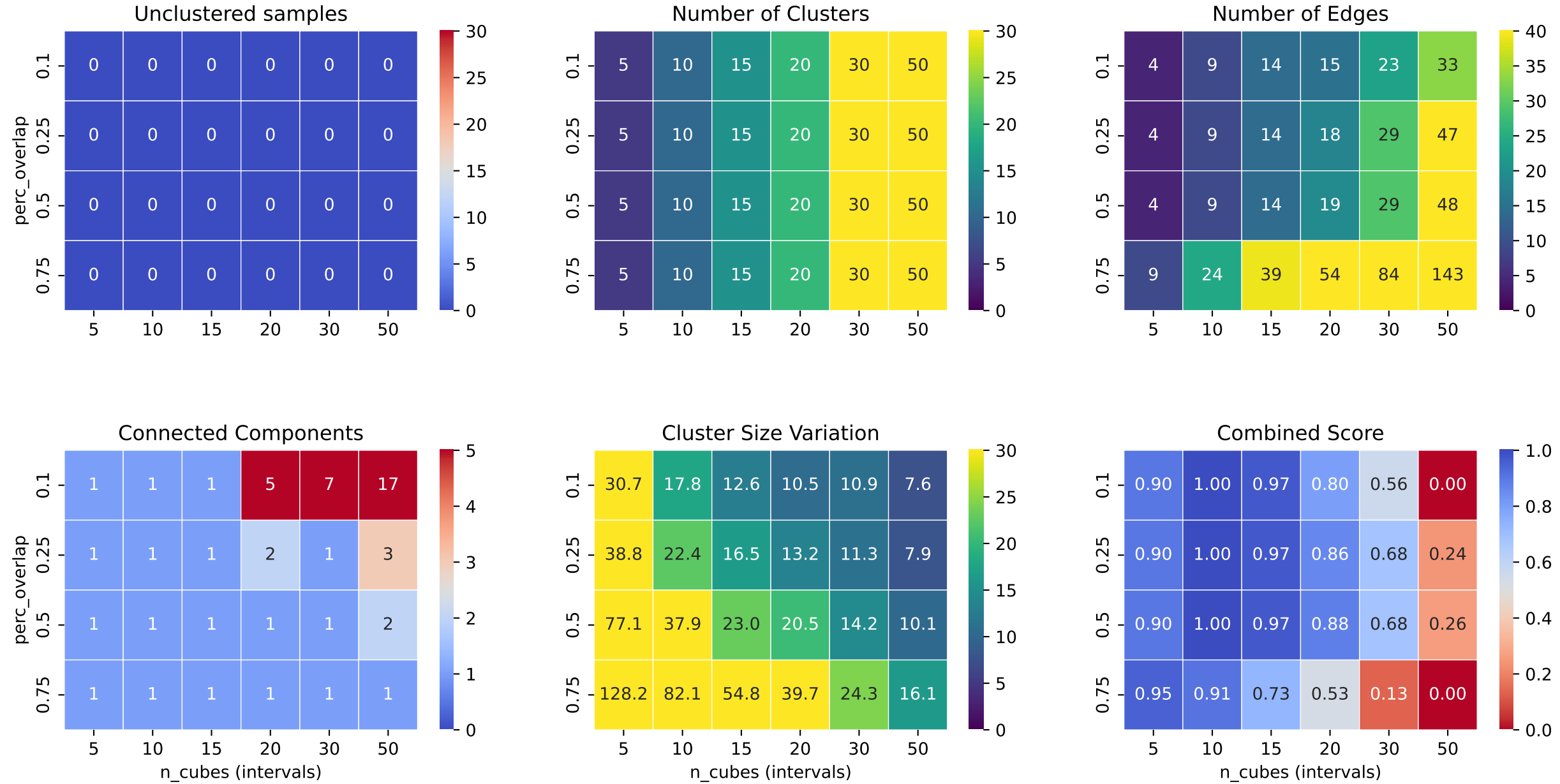
Projection: t-SNE (n_components=1)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=30.0)



Projection: t-SNE (n_components=1)

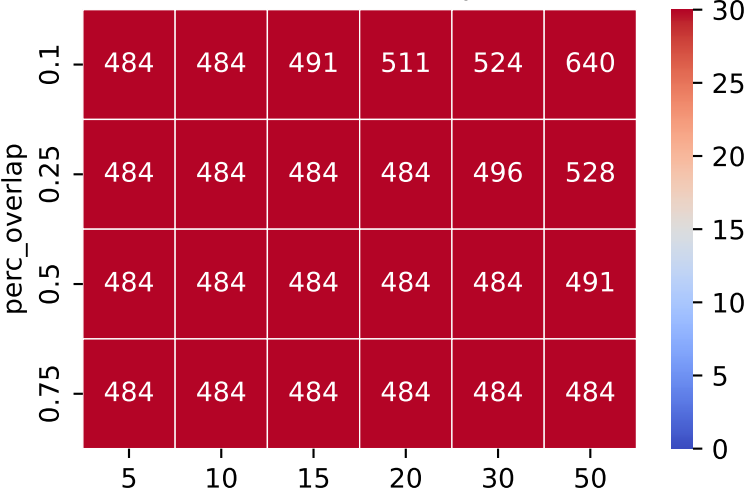
Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)



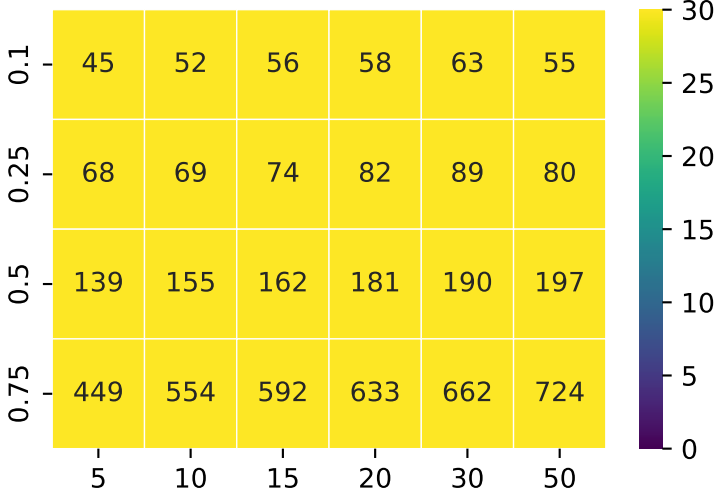
Projection: t-SNE (n_components=2)

Clustering: DBSCAN (eps=1.0)

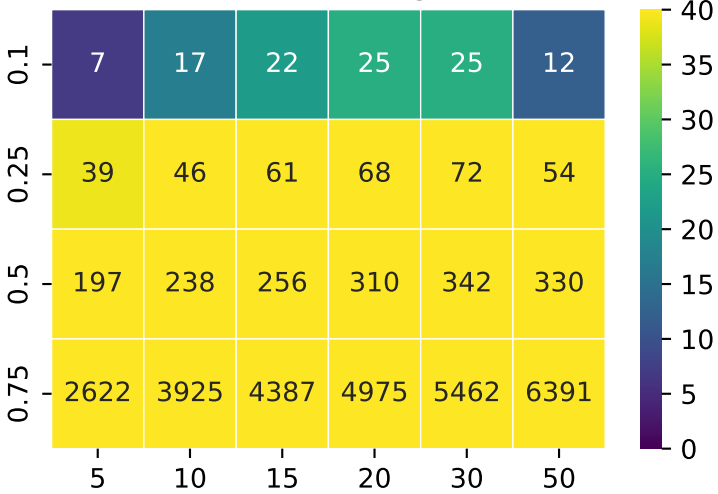
Unclustered samples



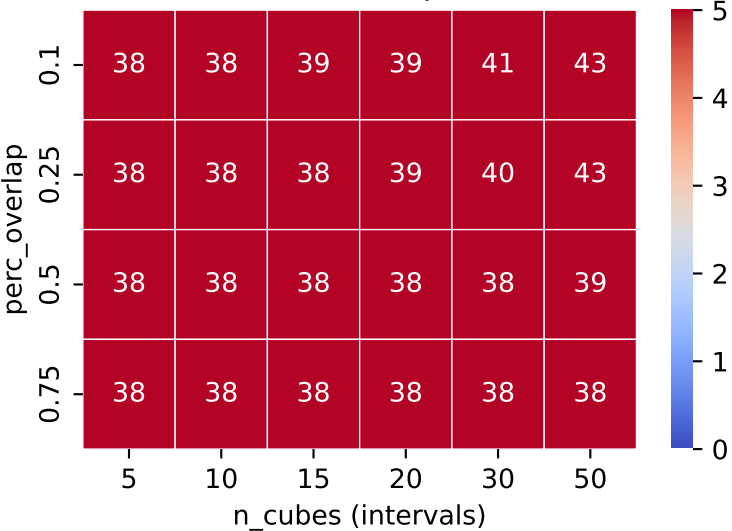
Number of Clusters



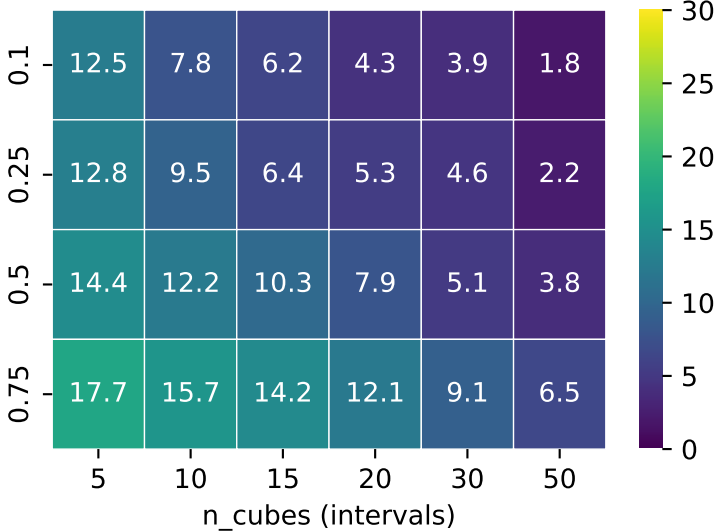
Number of Edges



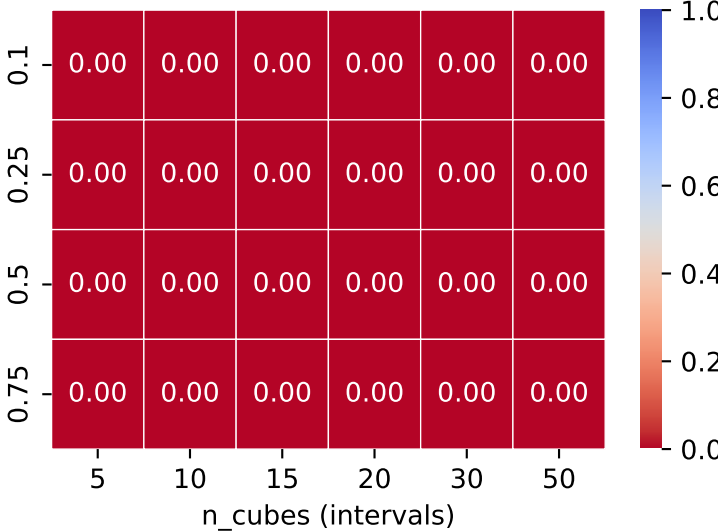
Connected Components



Cluster Size Variation



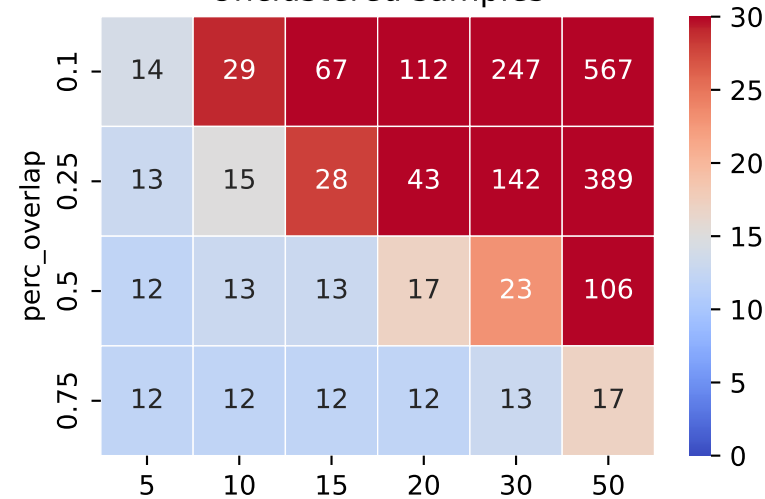
Combined Score



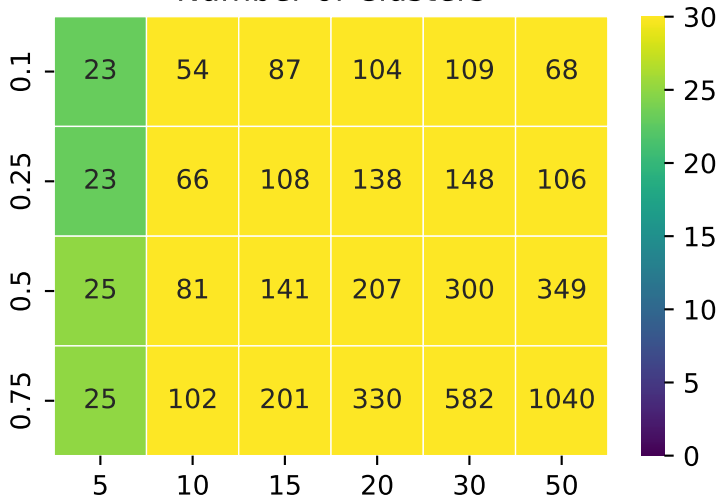
Projection: t-SNE (n_components=2)

Clustering: DBSCAN (eps=3.0)

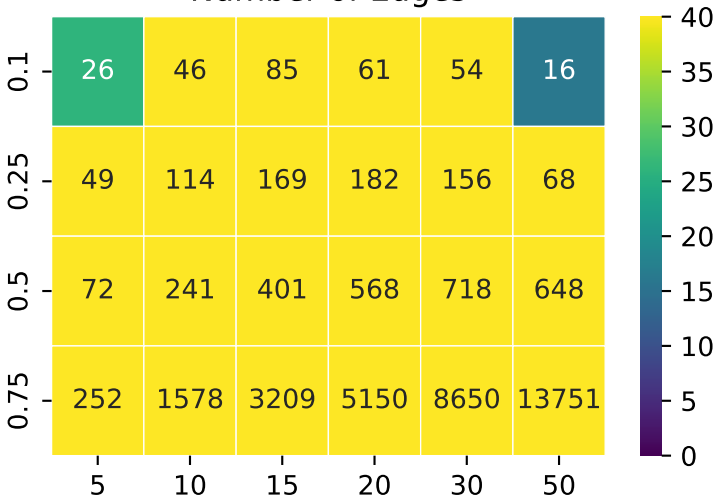
Unclustered samples



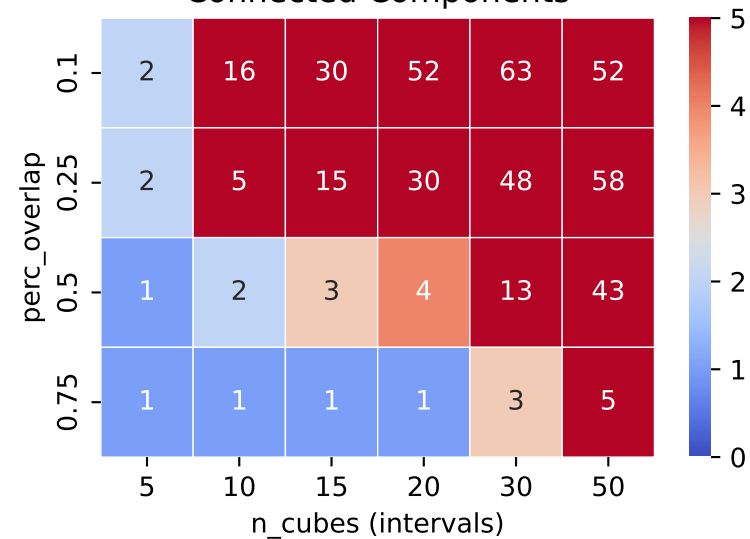
Number of Clusters



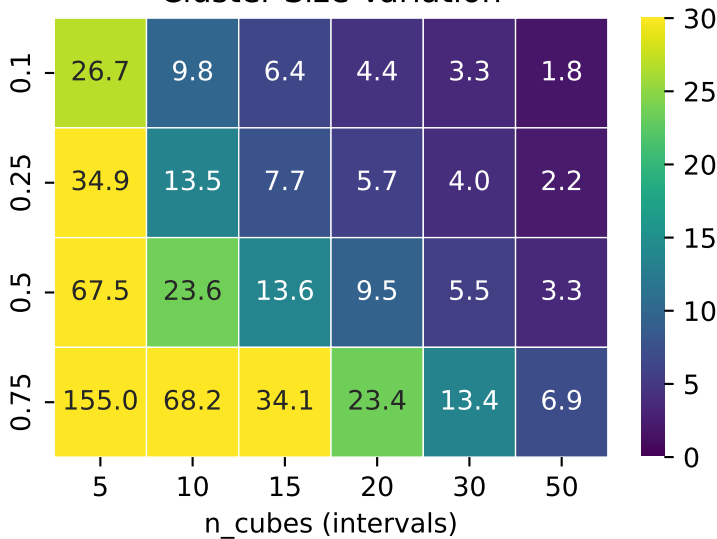
Number of Edges



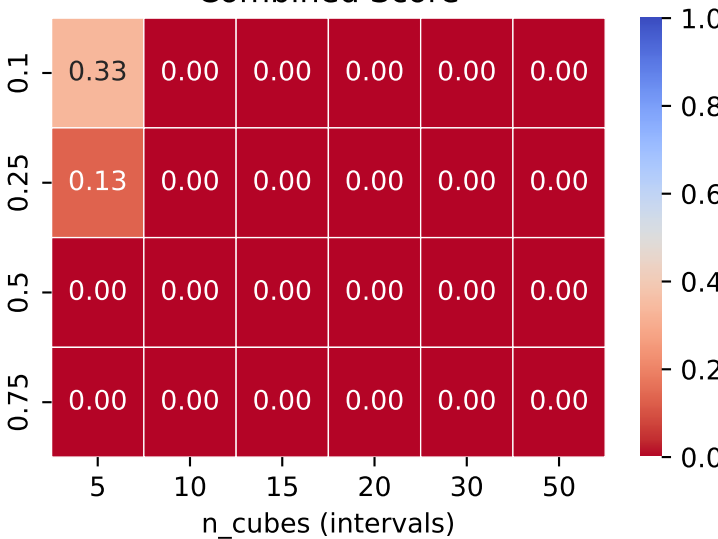
Connected Components



Cluster Size Variation



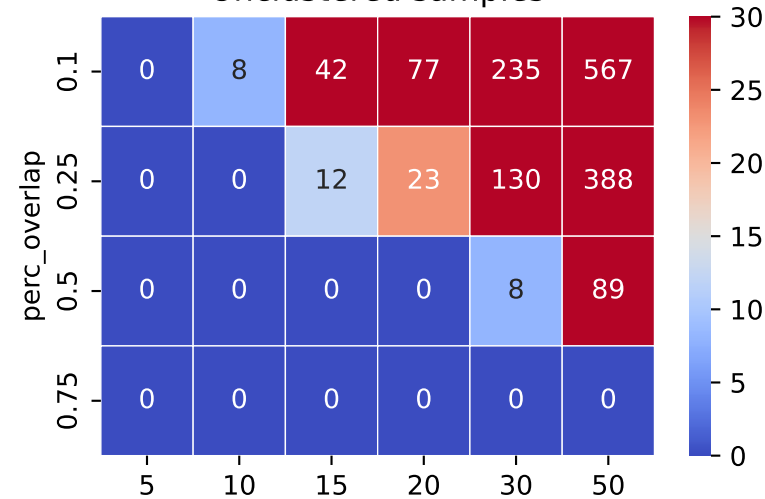
Combined Score



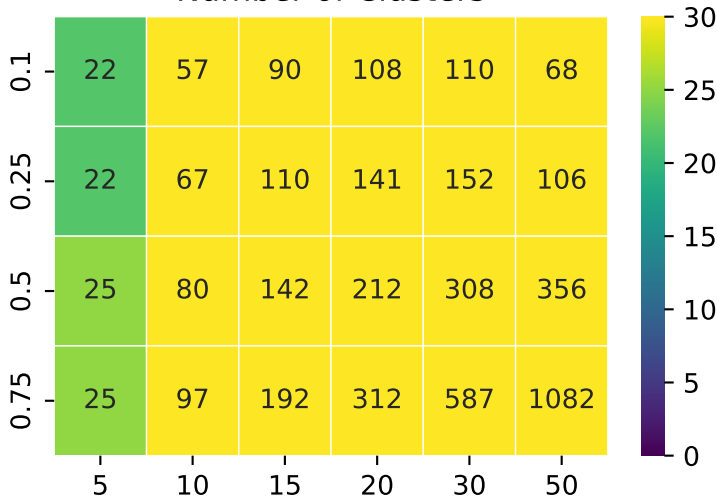
Projection: t-SNE (n_components=2)

Clustering: DBSCAN (eps=5.0)

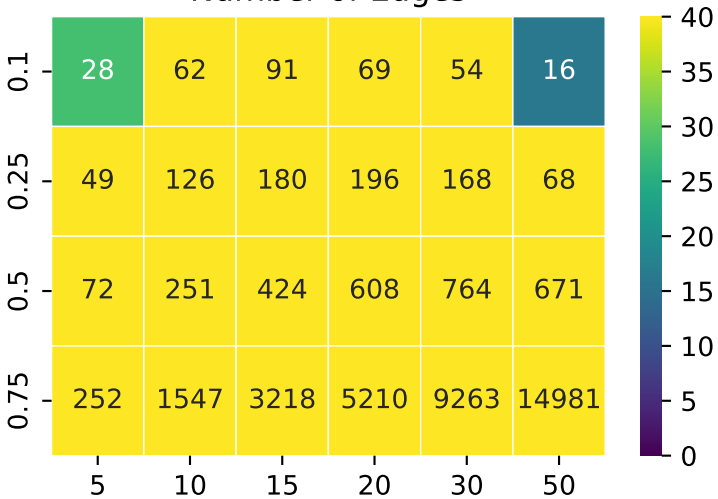
Unclustered samples



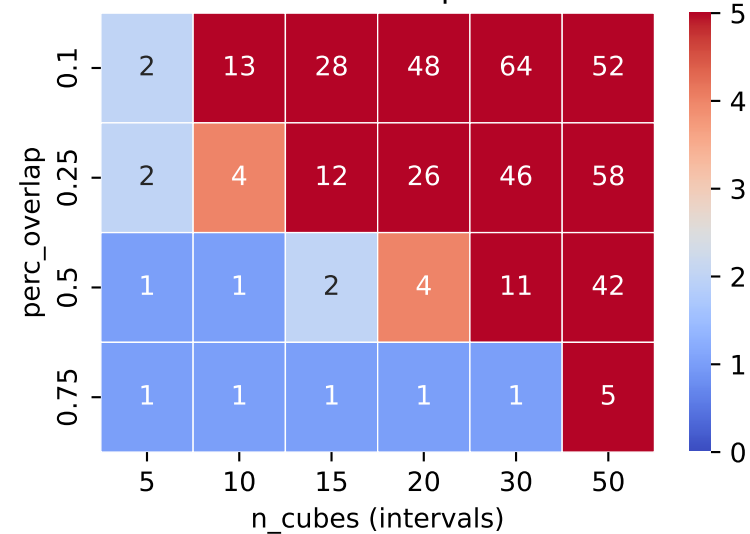
Number of Clusters



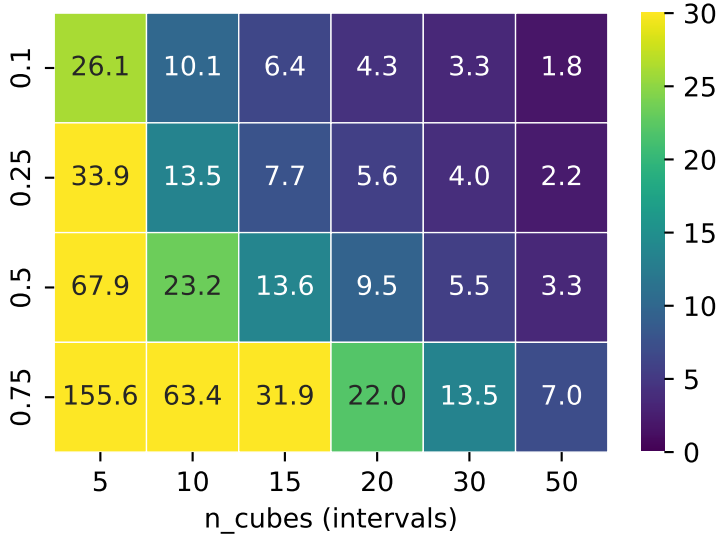
Number of Edges



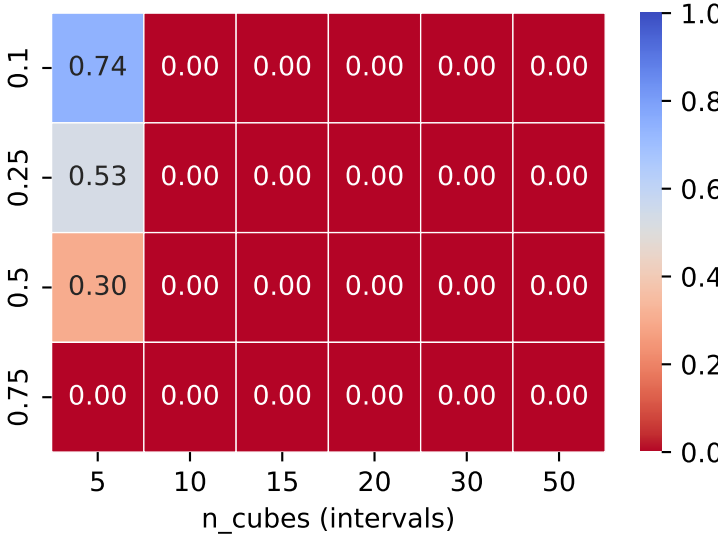
Connected Components



Cluster Size Variation



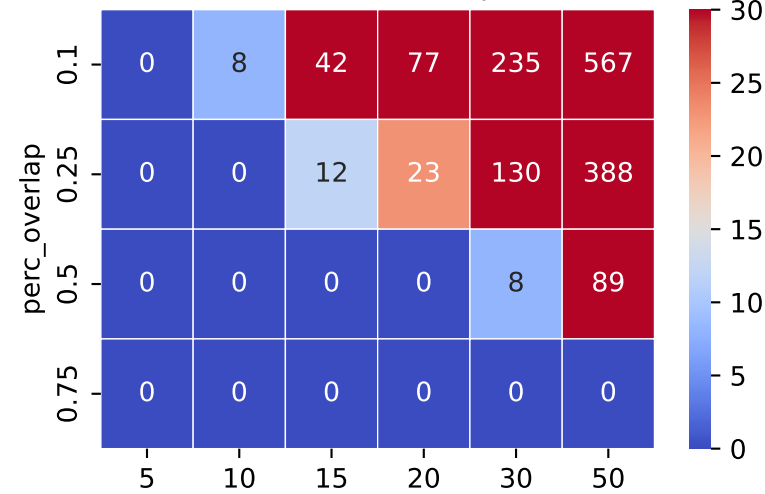
Combined Score



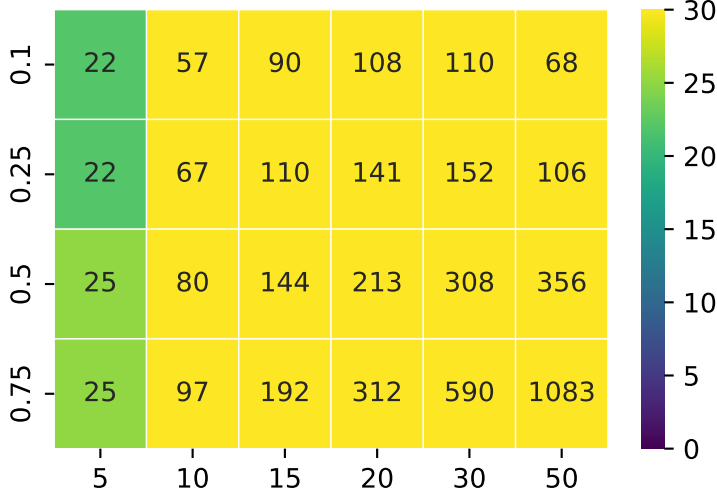
Projection: t-SNE (n_components=2)

Clustering: DBSCAN (eps=10.0)

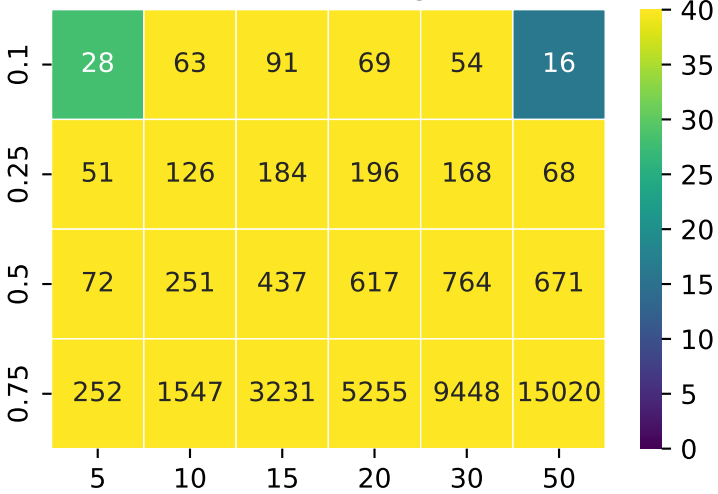
Unclustered samples



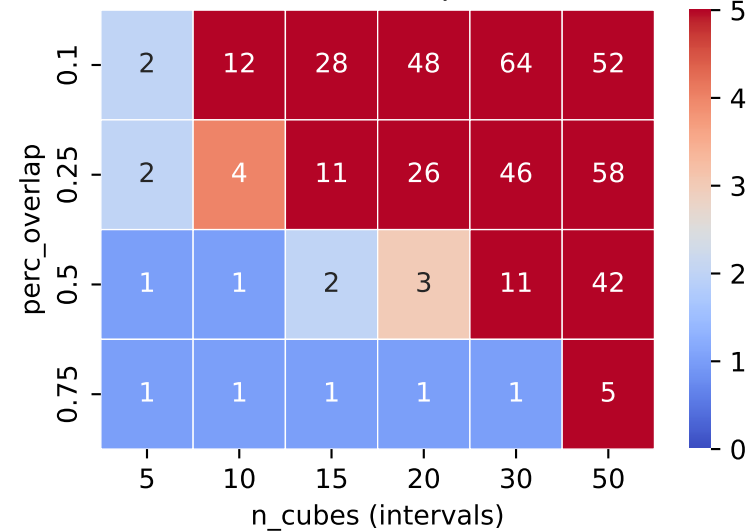
Number of Clusters



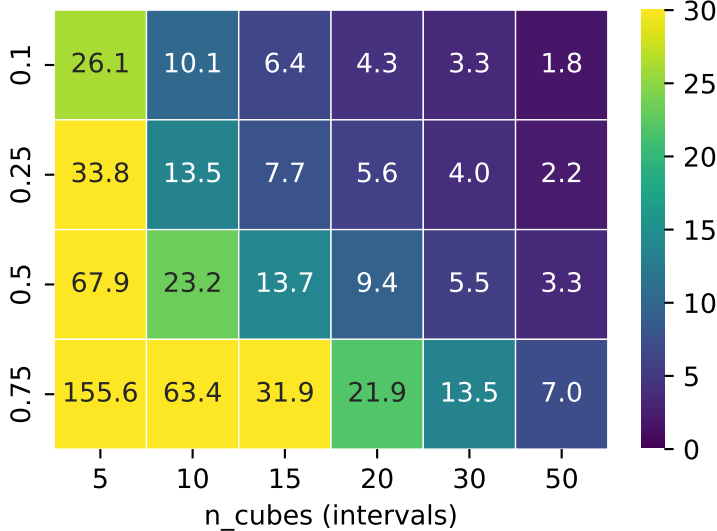
Number of Edges



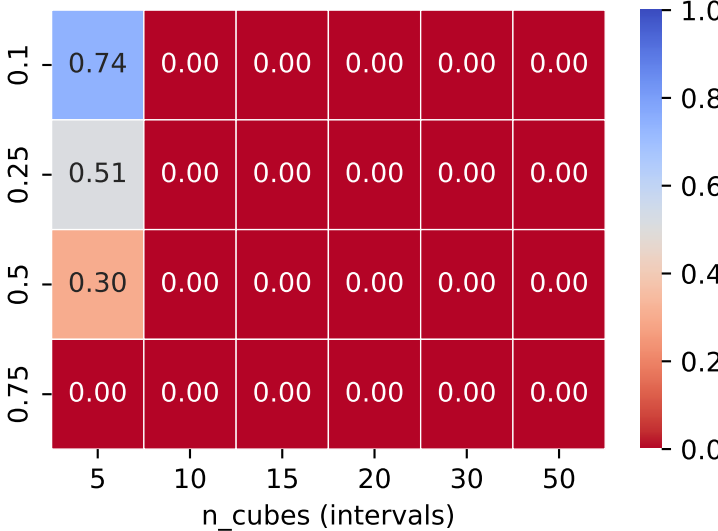
Connected Components



Cluster Size Variation



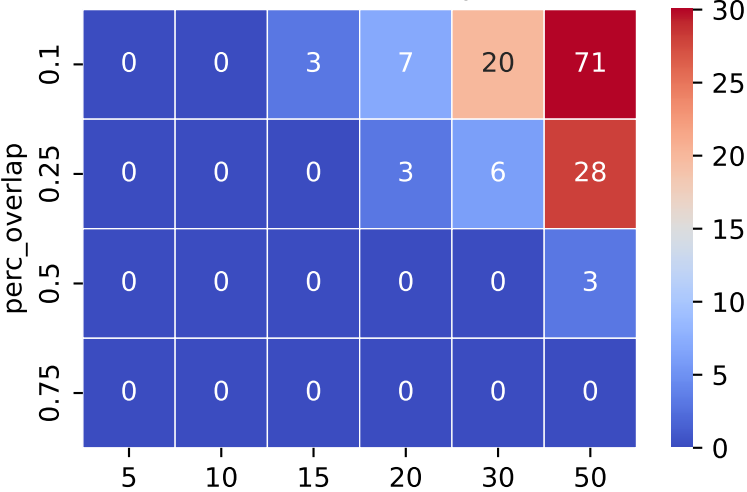
Combined Score



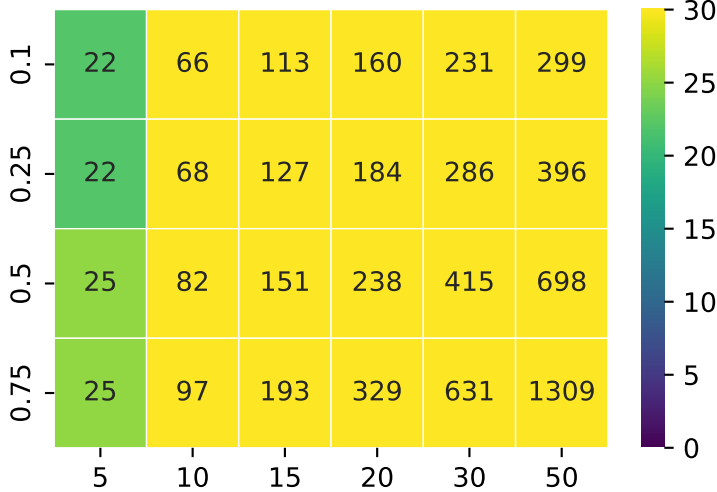
Projection: t-SNE (n_components=2)

Clustering: Bayesian Gaussian Mixture (n_components=1)

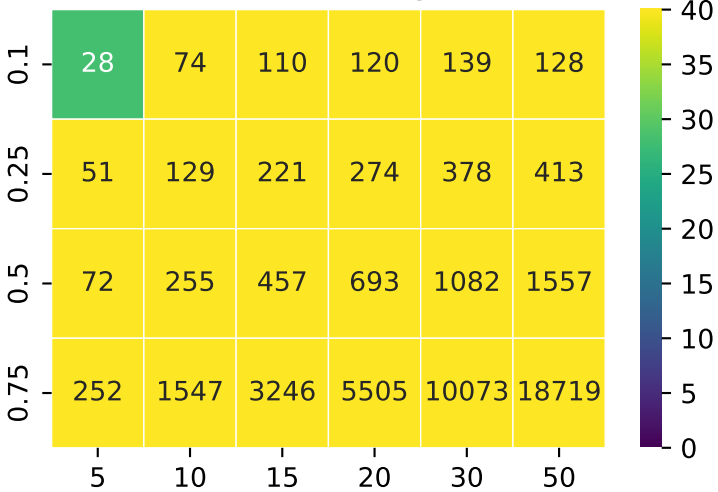
Unclustered samples



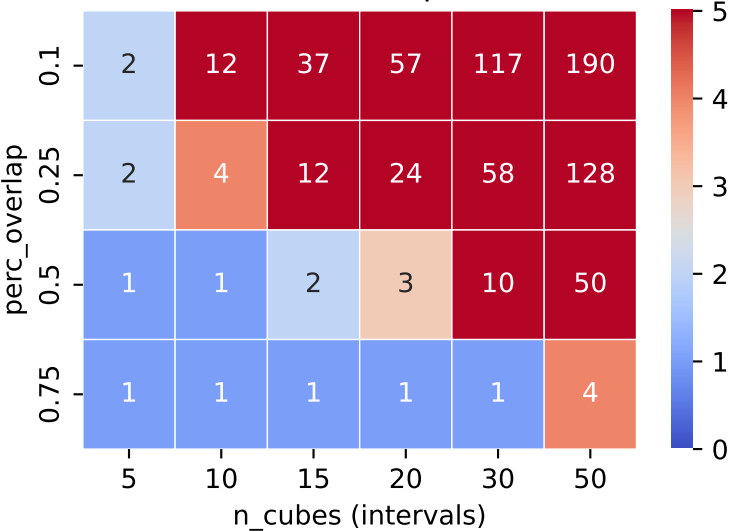
Number of Clusters



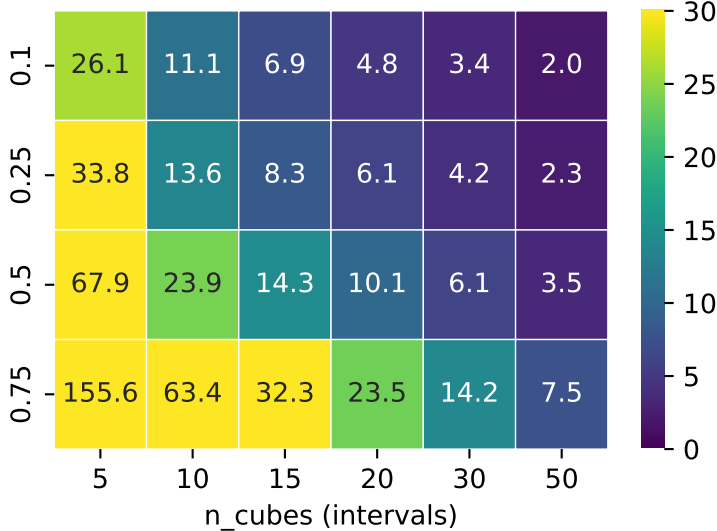
Number of Edges



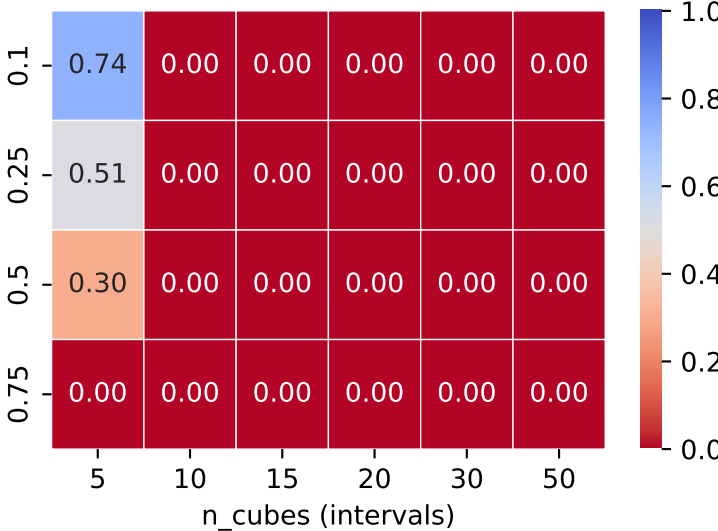
Connected Components



Cluster Size Variation

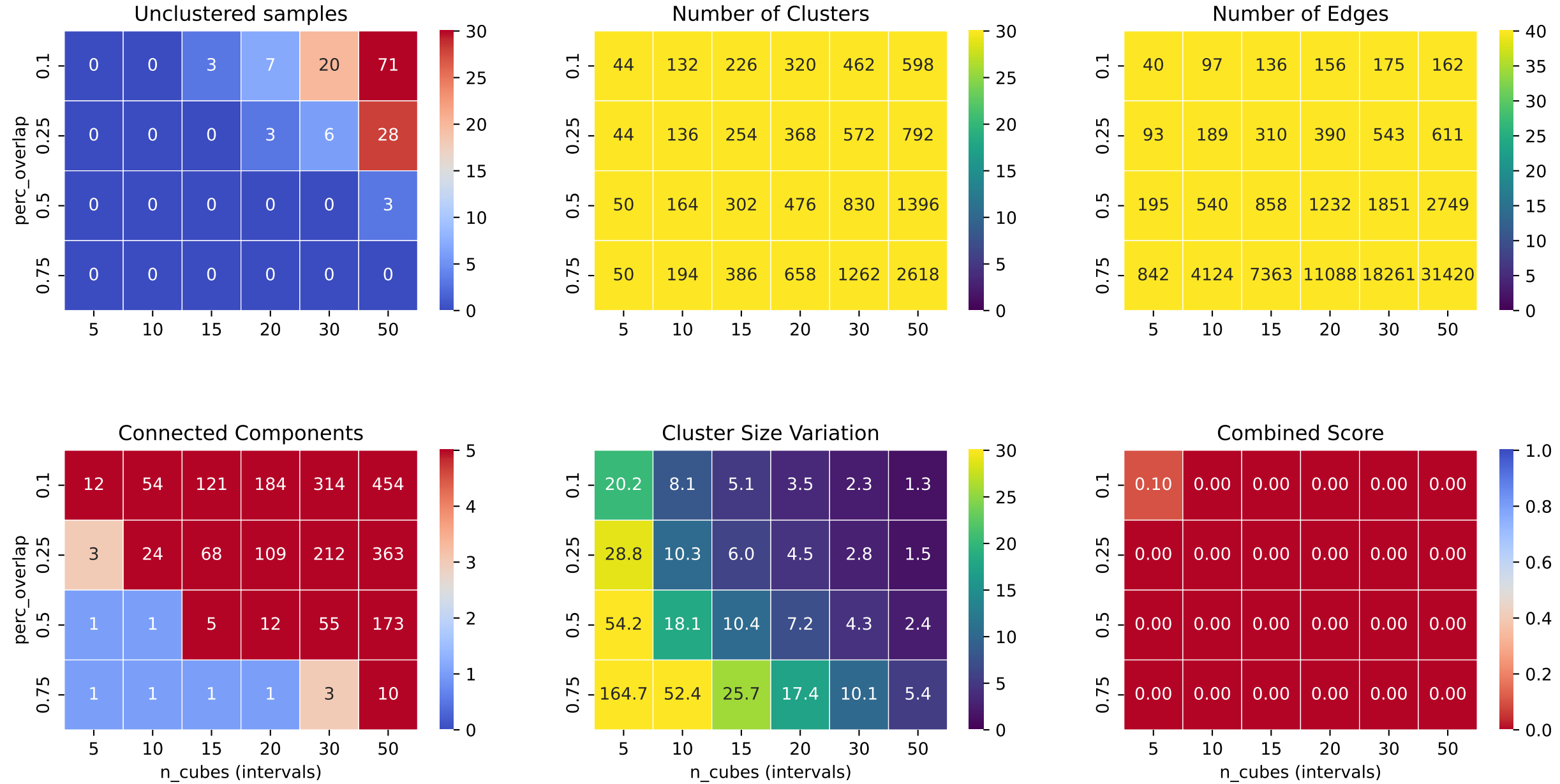


Combined Score



Projection: t-SNE (n_components=2)

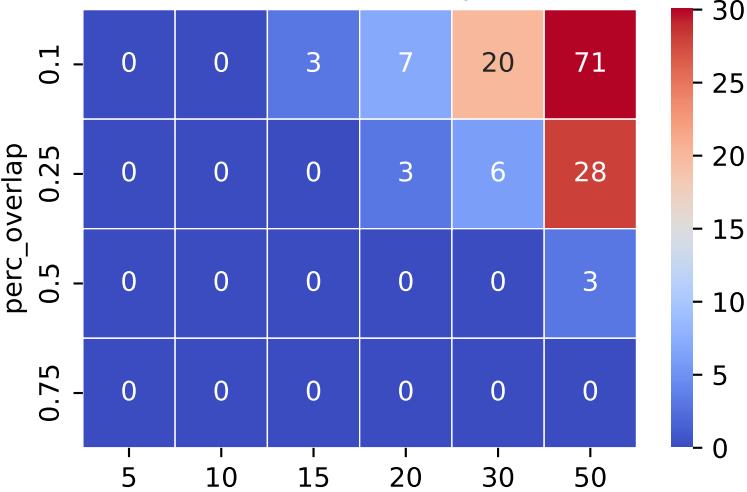
Clustering: Bayesian Gaussian Mixture (n_components=2)



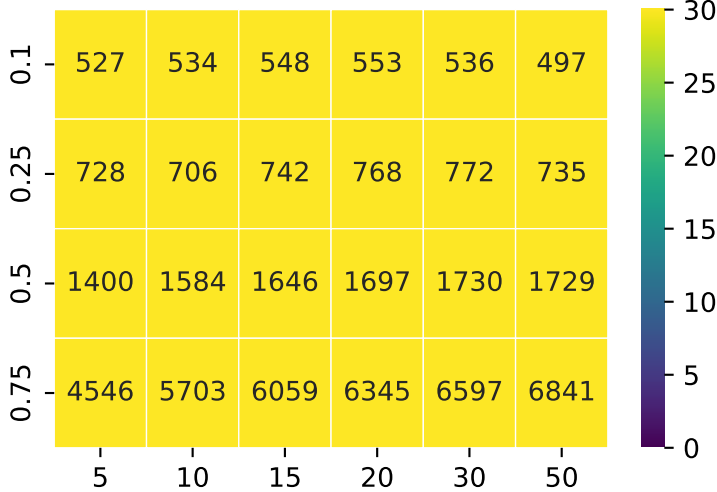
Projection: t-SNE (n_components=2)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=1.0)

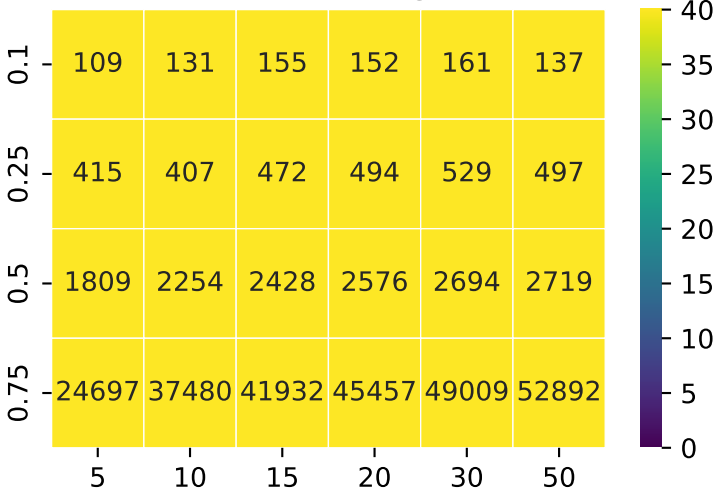
Unclustered samples



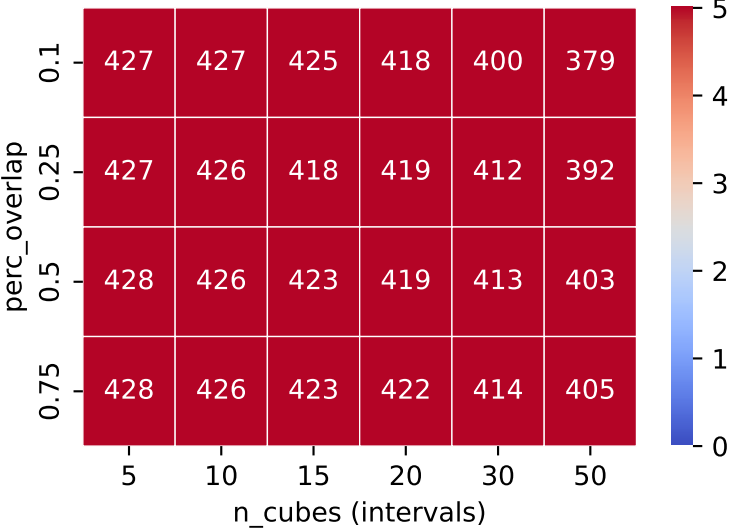
Number of Clusters



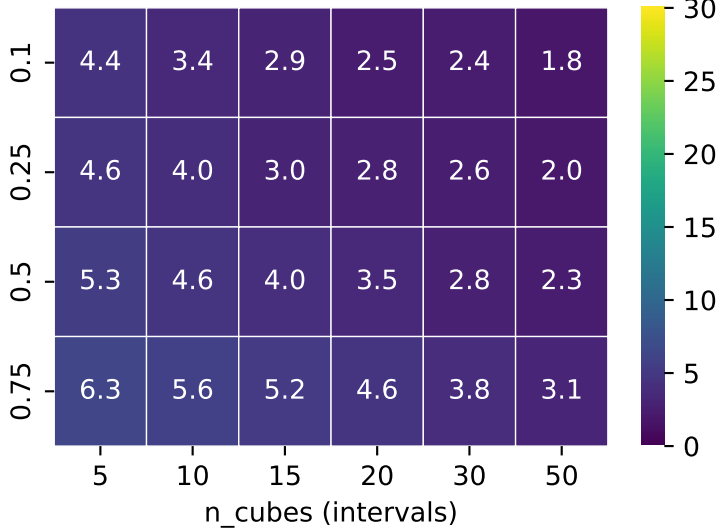
Number of Edges



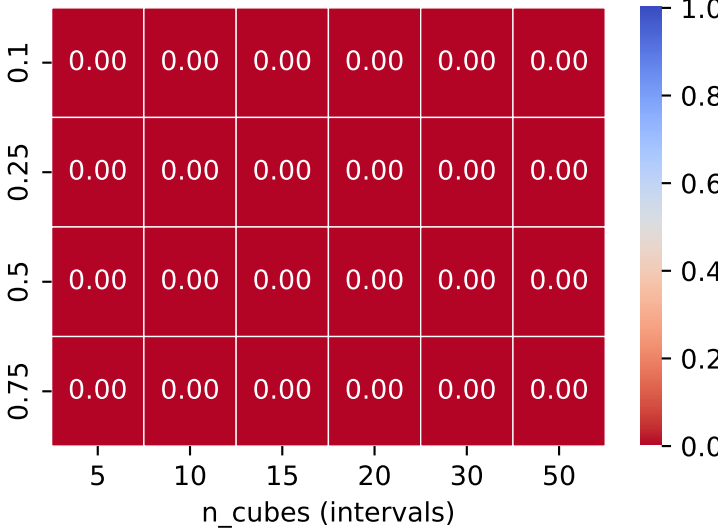
Connected Components



Cluster Size Variation

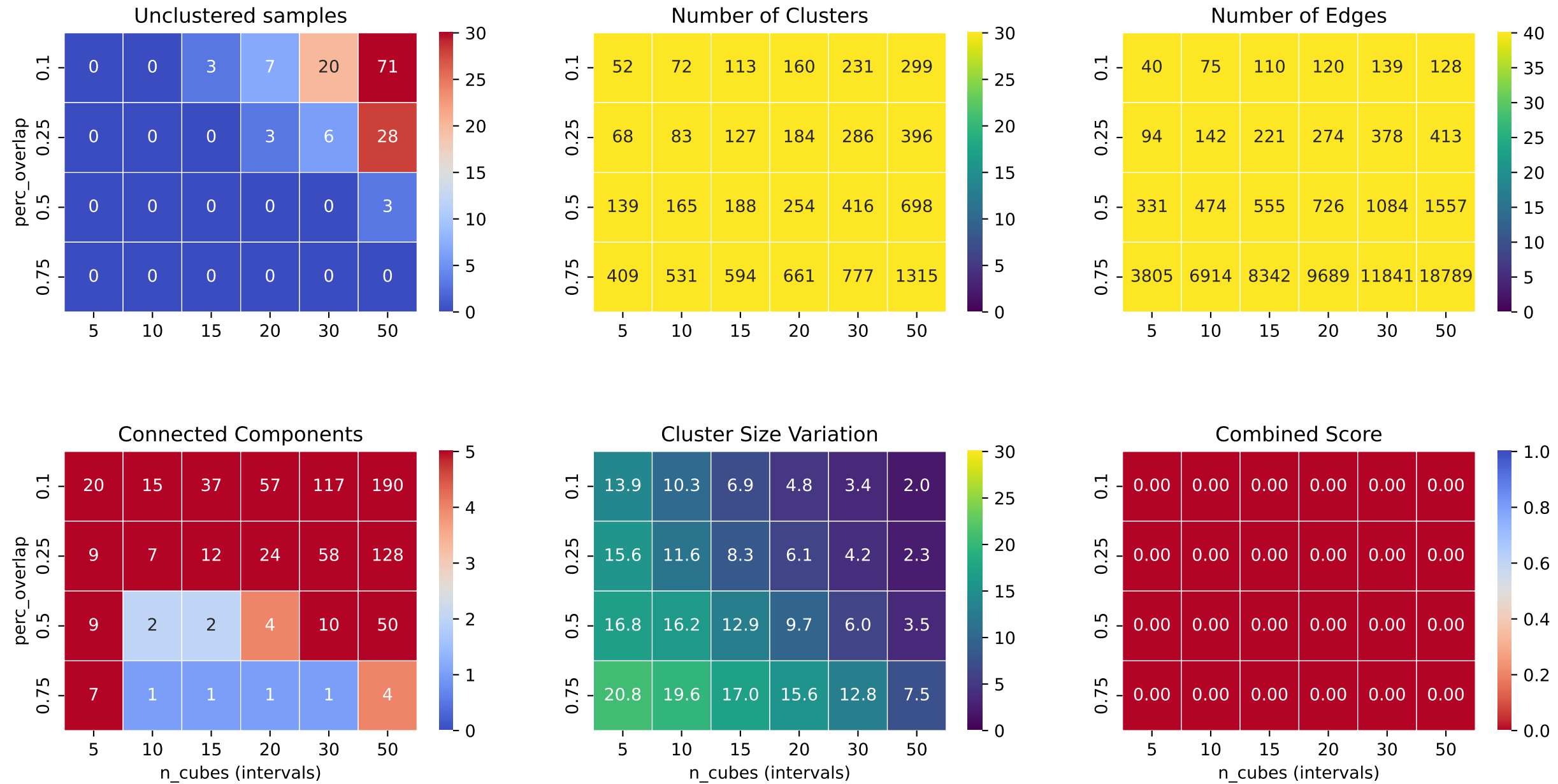


Combined Score



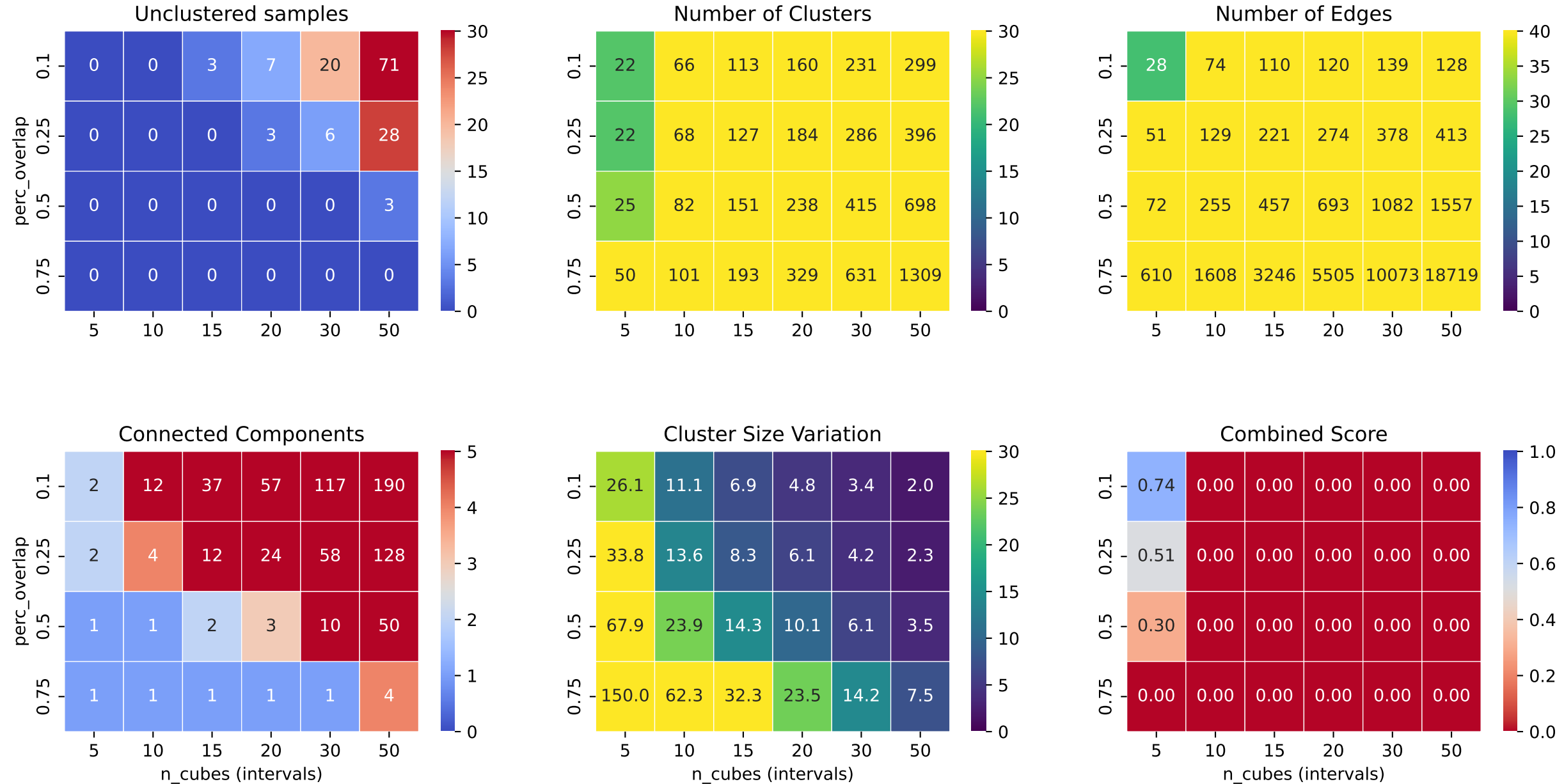
Projection: t-SNE (n_components=2)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=10.0)



Projection: t-SNE (n_components=2)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=30.0)



Projection: t-SNE (n_components=2)

Clustering: Hierarchical Agglomerative Clustering (distance_threshold=50.0)

